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**CONVERSATIONAL HIGH-RESOLUTION
MASS SPECTROGRAPHIC DATA REDUCTION**

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16. Abstract <p>A FORTRAN IV program is described which reduces the data obtained from a high-resolution mass spectrograph. The program (1) calculates an accurate mass for each line on the photoplate, and (2) assigns elemental compositions to each accurate mass. The program is intended for use in a time-shared computing environment and makes use of the conversational aspects of time-sharing operating systems.</p>					
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CONVERSATIONAL HIGH-RESOLUTION MASS SPECTROGRAPHIC

DATA REDUCTION

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SUMMARY

A FORTRAN IV program is described which reduces the data obtained from a high-resolution mass spectrograph. The program (1) calculates an accurate mass for each line on the photoplate, and (2) assigns elemental compositions to each accurate mass. The program is intended for use in a time-shared computing environment and makes use of the conversational aspects of time-sharing operating systems.

INTRODUCTION

Since the organic chemist first realized the potential of high-resolution mass spectrometry as an analytical tool (ref. 1), a great effort has been spent in developing computer software to process the large amounts of data generated (refs. 2, 3, and 4). The first programs were written for use on large batch computer systems. Recently, however, the emphasis has shifted to programing for relatively small, dedicated computer systems, with the data acquisition and reduction in real-time (refs. 5 and 6).

Obviously, a real-time data system has many advantages, the best of which is immediate data analysis and presentation. In the off-line schemes, typically a day or more separated the data acquisition and the computer analysis of the mass spectra. A disadvantage of the real-time systems is that it takes approximately 20 seconds (ref. 6) (for resolution of one part in 10,000) to acquire a mass spectrum. If greater resolution than one part in 10,000 is desired, even more time is necessary to obtain the spectrum. For certain applications, for example, a directly coupled gas-chromatograph mass spectrometer, the sample concentration may be so small that it does not remain in the mass spectrometer long enough for the complete mass spectrum to be acquired. For these kinds of applications a photodetection system, which is an integrating detector, is superior to the real-time system.

In the mass spectrometers used in real-time systems (ref. 7), the beam of ions comes to focus at a point. The magnetic field or ion accelerating voltage of the mass spectrometer must be scanned to sweep all the ion types past an electron multiplier at the point of focus. For samples of small concentration, as mentioned above, there may not be enough time to complete the scanning process. On the other hand, in a mass spectrograph (ref. 7), the ion optics of the instrument bring the sample beam to focus in a plane, and a photoplate placed in this plane can continuously record all the ions of the entire mass spectrum. With this scheme, even for small sample quantities, a complete mass spectrum can be obtained. For most of the analytical problems encountered in this laboratory, a photoplate recording system is the preferred method of data acquisition.

The computer program described below is designed to reduce data acquired off-line from a photoplate, but, because it runs on a time-sharing computer, it has some of the advantages of a real-time program. A day or more still separates acquisition of the mass spectrum and its computer analysis, but after the program is in use, it communicates with the user at a conversational terminal and accepts various inputs from the user during execution. As will be discussed later, parameters that affect the calculations in the program must be selected on the basis of various characteristics of each photoplate. With this conversational program, the selections can be made and changed while the program is running, the same as in a real-time system.

SYSTEM OVERVIEW

The high-resolution mass spectrograph (CEC 21-110B) employed at Ames Research Center is of the Mattuch-Herzog geometry. The sample to be analyzed is introduced into the ion source chamber of the mass spectrograph, and is ionized and fragmented by a collimated beam of thermally emitted 70 volt electrons. The positively charged fragments of the sample are accelerated through a focusing slit and passed into a magnetic field where ion-type groupings separate according to their mass. Each charged fragment type forms a slit image on a photoplate placed in the focal plane of the instrument. The photoplate can record up to 45 exposures (mass spectra) of typically several hundred slit images (peaks) each.

Each mass spectrum is transcribed from the photoplate to seven track BCD magnetic tape by a custom-designed comparator-microphotometer interfaced with a tape writing unit. The exposed photographic plate is scanned past the optical density measuring photomultiplier on a precision driven table. When the circuitry senses a line on the photoplate, the table is slowed to tape writing speed, and optical transmittance data are passed to the digitizing circuitry every one-half micron. After the last optical density reading in a slit image profile is recorded, the distance of this last reading, in half microns from the low mass end of the photoplate, is encoded.

At the conclusion of one complete spectrum transcription, the data tape will contain a profile and a distance value for each peak in the mass spectrum. This mass spectrograph data tape is read into disk storage on the Ames Research Center, IBM 360 Model 67 computer under an IBM supplied time-sharing operating system (TSS) and becomes the input file for the data reduction program.

From the shape of the line profiles and the distances of the last optical density readings in each profile, the position of the center of each peak can be determined to within an error of less than one-half micron. The masses of each ion type are related to each other by the square of the distance that separates them. Peaks of known mass are interspersed in the sample spectra by having a mass standard (perfluoroalkane) always present in the ionizing chamber. These peaks of known mass are used to calculate a distance to mass relationship for the length of the photoplate. From this relationship, the masses of the ion types present in the sample spectrum can be determined to within an error of a few parts per million. This mass measurement accuracy is, for the vast majority of problems, sufficient to assign unambiguous elemental compositions to each fragment in the sample spectrum. The final output is a tabular listing of the accurate mass and elemental composition of each peak in the mass spectrum.

COMPUTER PROGRAM DESCRIPTION

The computer program can be logically divided into three sections: peak center calculation, standard mass identification, and accurate mass calculation and element fitting.

Peak Center Calculation

Because the accurate mass calculated for each peak in the mass spectrum can only be determined within the error of the peak center calculations, the determination of the center of each peak profile is the most important aspect of the data reduction process. For an ideal line profile, whose shape is a gaussian distribution curve, the accurate center is easily discerned. In general, however, not all profiles in a spectrum will have a gaussian shape. There will be profiles with complicated shapes that describe lines which are not completely separated, and there will be wide profiles with flat tops that correspond to overexposed lines on the photographic plate. In addition, always superimposed on each profile will be a certain amount of random error commonly referred to as noise. An algorithm for peak center calculation must first attempt to filter out mathematically as much random error as possible, without degrading the underlying profile data, before addressing itself to the determination of the center of each peak.

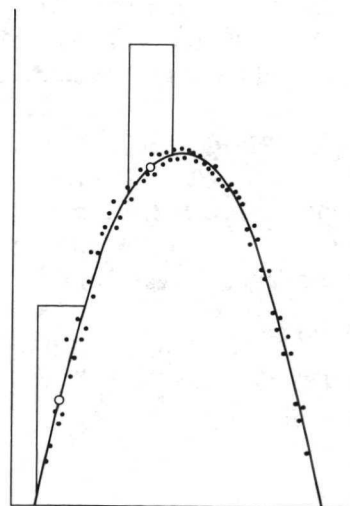
Data smoothing— On examining a plot of a peak profile, one would tend to draw through the points, a line which best fits them, thereby obtaining the best smooth curve through the points. Numerically, this can be done by fitting a least squares polynomial to the data profile.

The set of points to be smoothed is fit to some polynomial curve.

$$P(x) = \sum_{j=0}^N a_j x^j$$

The coefficients designated a are computed so that when the x values are substituted into the polynomial equation, the square of the differences between the computed values and the profile data points are a minimum. Methods used to calculate the a coefficients are described in most textbooks on numerical analysis (refs. 8 and 9). The degree of the polynomial (which can be varied from 1 to 10) and the number of points used (2 to 200) in the moving polynomial smoothing function can be selected and changed while the program is executing. The sketch shows how the smoothing process works, for an overlapping 11 point polynomial.

As in the first grouping in the sketch a new value for the central position in the group is calculated from the least squares polynomial. The point at the left is dropped and a new point added from the right. Again a central point is calculated from the new polynomial and the process is repeated until a smoothed curve is obtained for the entire profile. An intermediate grouping is also indicated in the sketch. All peak profiles in the mass spectrum are smoothed in this fashion.



Peak center calculation— The first derivative of each peak profile is calculated using overlapping polynomials as above. From the shape of the first derivative curve, the number of peaks and the center of each peak in the profile can be determined (ref. 10).

Not all line profiles are analyzed in this fashion. Narrow profiles which can only contain a single peak are analyzed by a scheme that uses less computer time. If the total number of points in the peak profile is below some minimum number (selected during the program execution), the profile is best fit to a gaussian curve and the center and height determined directly from this curve (ref. 4).

$$y = y_{\max} \exp \left[-\frac{5.545(x - x_{\max})^2}{2w^2} \right]$$

$$\ln y = \ln y_{\max} - \frac{5.545(x - x_{\max})^2}{2w^2}$$

$$\ln y = \left(\ln y_{\max} - \frac{5.545}{2w^2} x_{\max}^2 \right) + \left(\frac{5.545x_{\max}}{w^2} \right) x - \left(\frac{5.545}{2w^2} \right) x^2$$

- y intensity at position x
- y_{\max} intensity at the maximum
- x_{\max} position of the maximum
- w half-width

This second degree polynomial can be fit to the entire profile to yield directly the center and height of a peak.

Occasionally, a peak is so overexposed on the photoplate that its shape cannot be properly analyzed by the methods described above. For this contingency a provision exists whereby a profile with an excessively large number of data points (the number is selected during program execution) will be displayed for editing on the conversational terminal. Long leading edges and tails (which are a phenomenon caused by the emulsion on the photoplate during an overexposure) can be eliminated and the profile can otherwise be edited before any further analysis.

Peak intensity calculation— With electrical ion detection, the abundance of each ion type can be accurately determined, but the intensity measurement capabilities of a photoplate are very limited (ref. 11). Because of this limitation, no attempt is made at accurate ion abundance measurements. The largest intensity reading in each profile is used to calculate an approximate relative abundance for each ion type.

A list of peak centers, in millimeters from the left edge of the photoplate, and high intensity values for each peak in the mass spectrum are passed to the standard mass identification portion of the program.

Standard Mass Identification

As mentioned above, not all the lines in a spectrum are due to the analyzed sample; some arise from an internal mass standard (perfluoroalkane). It is the task of this section of the program to identify which peaks correspond to masses from the mass standard and to use these to establish a distance-to-mass relationship for the length of the photoplate.

Mass identification— To begin with, the distances of the first 40 peaks in the list are displayed on the conversational terminal. At this point the program must be supplied the known mass of at least two lines in the table. Using these two (or more) lines, the program will calculate distances where it expects to find 160 of the most prominent peaks in the perfluoroalkane spectrum. Each time the program is successful in its search, it incorporates the new line into its standard mass table. At the end of one search iteration, the program will use the table of identified masses to calculate a distance-to-mass relationship.

Distance to mass conversion— Theoretically, there is a linear relationship between the square root of the mass of each ion type and the distance that separates the ions (ref. 7). In practice, because of inhomogeneous areas in the magnetic field used for separation, the relationship is not exactly linear, but is best approximated by overlapping polynomial functions that are fitted to the standard mass values spread throughout the length of the photoplate. The degree of and the number of points spanned by the polynomials may be selected during the program execution. Generally, the best results are obtained from a second degree polynomial fit to four standard masses. The coefficients of the polynomial are calculated and saved for each overlapping set of calibration masses. To assess the accuracy of the conversion relationship, these polynomials are used to recalculate the masses of the identified standard mass lines, and this information is displayed at the conversational terminal. The difference between the calculated mass and the known mass of the lines of the calibration compound gives a very good idea of the accuracy of the distance-to-mass relationship in the area of interest. At this point misidentified peaks in the list may be deleted, and missed peaks may be added from the conversation terminal. If there has been any input from the user, the program loops back to the standard mass search routine and a new conversion relationship is calculated. This process may be repeated until one is sure that the calculated distance-to-mass relationship is correct.

Element fitting parameters— If there are no additions to or deletions from the standard mass list, the program prompts the user for element fitting parameters. A tolerance for element fitting must be chosen; it is the variance in millimass units between the calculated mass and the mass of an acceptable elemental composition. Next the elements considered for possible compositions must be selected. Carbon, hydrogen, nitrogen, and oxygen are always included in the candidate element table, and as many as seven additional elements may be added for consideration at execution time.

After the element fitting parameters are entered, the user may choose either (1) to write all information calculated thus far into a disk file, where it may be accessed by a program that uses the computer center line printer for output, or (2) to let the program continue with the final results displayed at the conversational terminal. Usually there will be too much information for listing on the relatively slow conversational terminal.

Mass Calculation and Element Fitting

Mass calculation— The distance-to-mass relationship just obtained, is used to calculate an accurate mass for all the lines in the mass spectrum. Each mass is calculated using the polynomial that was fit to the nearest group of calibration masses. Next, all masses corresponding to ion species commonly found in the background of the mass spectrometer and all ion types of the marker compound are eliminated from the list of calculated masses. It is this abbreviated list of accurate masses that represents the mass spectrum of the unknown sample.

Element fitting— All possible permutations of the elements submitted for element fitting are considered as compositions for each of the masses calculated. The minimum number of each isotope considered is zero. The maximum number varies as follows: for carbon-12 it is the integer resulting from the division of the accurate mass by 12; for carbon-13 it is one; for hydrogen it is twice the maximum number of carbon atoms plus eight; for nitrogen-14 it is two; for oxygen-16 it is three. All other isotopes are entered during program execution and the maximum number of each is one of the input parameters. All permutations whose masses fall within the millimass unit tolerance specified are displayed along with the calculated accurate masses.

Output— The final output consists of a relisting of the standard mass table of part II and a table of all calculated masses and their associated elemental compositions. If no permutations of elements fall within the tolerance, the calculated mass is listed with an appropriate comment. There is no provision for special handling of multiply-charged ions in the mass spectrum. Generally, if an obviously multiply-charged species (calculated to be in the vicinity of half an integral mass unit) is doubled, the corresponding singly-charged ion can be found at this value. The accurate mass of the multiply charged ion species is listed with no associated elemental compositions.

CONCLUSIONS

A typical standard mass listing is shown in appendix A. The largest deviation from the exact mass in the entire list is less than five parts per million. This accuracy compares favorably with previously reported high resolution data analysis programs. In addition, for directly coupled gas-chromatograph mass spectrometry applications, a photoplate detection system used with the conversational data reduction system described here is more sensitive and more efficient than previously reported systems.

Ames Research Center

National Aeronautics and Space Administration

Moffett Field, Calif. 94035, March 22, 1973.

APPENDIX A

SAMPLE OF STANDARD MASS ACCURACY

	DISTANCE	CALCULATED MASS	EXACT MASS	DIFFERENCE IN MMU	DIFFERENCE IN PPM
1	15.3497	26.01569	26.01565	0.04	1.47
2	16.9022	27.02342	27.02347	-0.06	2.04
3	18.3888	28.00616	28.00615	0.01	0.29
4	22.7609	30.99844	30.99840	0.04	1.34
5	24.1629	31.98985	31.98983	0.02	0.51
6	29.5927	35.97659	35.97668	-0.09	2.44
7	34.7268	39.96239	39.96239	0.00	0.02
8	39.6593	43.98983	43.98983	-0.00	0.02
9	46.6147	49.99686	49.99680	0.06	1.10
10	47.7399	51.00465	51.00463	0.02	0.43
11	52.0936	54.99833	54.99840	-0.07	1.19
12	59.3566	61.99684	61.99680	0.04	0.71
13	67.1731	69.99841	69.99856	-0.15	2.20
14	70.9115	73.99688	73.99680	0.08	1.10
15	77.2184	80.99519	80.99521	-0.02	0.25
16	81.5609	85.99656	85.99681	-0.25	2.90
17	87.4290	92.99544	92.99521	0.23	2.52
18	93.0803	99.99501	99.99521	-0.20	2.00
19	96.9997	104.99532	104.99521	0.11	1.04
20	102.3302	111.99335	111.99361	-0.26	2.33
21	106.0392	116.99518	116.99521	-0.03	0.26
22	111.0970	123.99365	123.99361	0.04	0.34
23	116.0157	130.99141	130.99202	-0.61	4.68
24	116.7109	131.99539	131.99537	0.02	0.18
25	122.8240	140.99512	140.99521	-0.09	0.61
26	124.1546	142.99197	142.99201	-0.04	0.28
27	126.7720	146.96262	146.96248	0.14	0.95
28	128.7443	149.99033	149.99041	-0.08	0.54
29	131.9597	154.99195	154.99201	-0.06	0.41
30	136.3722	161.99051	161.99041	0.10	0.62
31	139.4679	166.99177	166.99201	-0.24	1.46
32	140.6919	168.98871	168.98882	-0.11	0.62
33	143.7246	173.99051	173.99041	0.10	0.59
34	146.7139	178.99164	178.99201	-0.37	2.08
35	146.8967	180.98895	180.98882	0.13	0.70
36	150.8293	185.99030	185.99041	-0.11	0.61
37	154.8685	192.98889	192.98882	0.07	0.37
38	158.8352	199.98709	199.98722	-0.13	0.65
39	161.6289	204.98894	204.98882	0.12	0.58
40	165.4817	211.98724	211.98722	0.02	0.08
41	168.1976	216.98875	216.98882	-0.07	0.30
42	171.9456	223.98732	223.98722	0.10	0.45
43	174.5880	228.98836	228.98882	-0.46	2.01
44	175.6358	230.98565	230.98562	0.03	0.14
45	178.2386	235.98724	235.98722	0.02	0.10
46	180.8154	240.98862	240.98882	-0.20	0.82
47	181.8369	242.98585	242.98562	0.23	0.95
48	184.3764	247.98717	247.98722	-0.05	0.22
49	187.8883	254.98563	254.98562	0.01	0.04
50	190.3679	259.98745	259.98722	0.23	0.90
51	191.3517	261.98396	261.98402	-0.06	0.24
52	193.8005	266.98578	266.98562	0.16	0.59
53	194.7711	268.98230	268.98242	-0.12	0.45
54	197.1875	273.98413	273.98402	0.11	0.41
55	199.5817	278.98529	278.98562	-0.33	1.17
56	200.5325	280.98243	280.98242	0.01	0.04
57	202.8978	285.98399	285.98402	-0.03	0.11
58	205.2427	290.98575	290.98562	0.13	0.46
59	206.1733	292.98258	292.98242	0.16	0.54
60	208.4914	297.98402	297.98402	-0.00	0.01
61	211.7035	304.98241	304.98242	-0.01	0.03
62	217.1272	316.98269	316.98242	0.27	0.86
63	218.0198	318.97910	318.97923	-0.13	0.41
64	220.2442	323.98096	323.98083	0.13	0.39
65	222.4508	328.98184	328.98242	-0.58	1.76
66	223.3280	330.97922	330.97923	-0.01	0.03
67	225.5114	335.98091	335.98083	0.08	0.25
68	227.6800	340.98186	340.98242	-0.56	1.65
69	228.5419	342.97977	342.97923	0.54	1.59
70	230.6881	347.98047	347.98083	-0.36	1.03
71	233.6681	354.97938	354.97922	0.16	0.44
72	235.7785	359.98078	359.98083	-0.05	0.13
73	238.7097	366.97938	366.97923	0.15	0.42
74	239.5406	368.97603	368.97603	0.00	0.01
75	243.6715	378.97910	378.97923	-0.13	0.34
76	244.4894	380.97594	380.97603	-0.09	0.24
77	246.5286	385.97791	385.97763	0.28	0.72
78	249.3603	392.97586	392.97603	-0.17	0.42
79	254.1604	404.97645	404.97603	0.42	1.05

	DISTANCE	CALCULATED MASS	EXACT MASS	DIFFERENCE IN MMU	DIFFERENCE IN PPM
80	257.3239	412.98206	412.98242	-0.36	0.86
81	258.8923	416.97604	416.97603	0.01	0.02
82	263.5553	428.97626	428.97603	0.23	0.54
83	264.3252	430.97260	430.97284	-0.24	0.55
84	266.2459	435.97467	435.97444	0.23	0.52
85	268.1541	440.97582	440.97603	-0.21	0.48
86	268.9139	442.97233	442.97284	-0.51	1.14
87	273.4419	454.97284	454.97284	0.00	0.01
88	277.9075	466.97284	466.97284	0.00	0.00
89	282.3171	478.97289	478.97284	0.05	0.10
90	283.0455	480.96973	480.96964	0.09	0.20
91	287.3917	492.96960	492.96964	-0.04	0.07
92	291.6849	504.96941	504.96964	-0.23	0.46
93	295.9264	516.96986	516.96964	0.22	0.43
94	300.1182	528.96730	528.96964	-2.34	4.43
95	300.8127	530.96465	530.96445	0.20	0.38
96	304.9507	542.96604	542.96645	-0.41	0.75
97	309.0428	554.96634	554.96645	-0.11	0.21
98	313.0897	566.96659	566.96645	0.14	0.25
99	317.0956	578.96471	578.96645	-1.74	3.01
100	317.7595	580.96341	580.96325	0.16	0.27
101	321.7162	592.96277	592.96325	-0.48	0.80
102	325.6343	604.96374	604.96325	0.49	0.80
103	329.5140	616.96308	616.96325	-0.17	0.27

APPENDIX B

PROGRAM LISTING

```

C   JANUARY 9,1973                               MICHAEL ROMIEZ
C
C   THIS PROGRAM WILL CONTROL THE READING OF THE AMES HIGH RESOLUTION DATA TAPE INTO A
C   DISK FILE WHICH CAN BE ACCESSED BY THE MAIN DATA REDUCTION PROGRAM.
C
C   THE DATA ON THE TAPE IS ENCODED IN SEVEN TRACK BCD FORMAT; IT IS CONVERTED INTO EBCDIC
C   BY AN AMES LIBRARY SUBROUTINE ALBYTES. THE DATA IS CONVERTED TO INTEGER FORMAT BY AN
C   ASSEMBLY SUBROUTINE TAPERD. THE RAW DATA IN INTGER FORMAT IS WRITTEN INTO A DISK FILE.
C
C                                     VARIABLES
C
C   IBYTES          - THE NUMBER OF BCD CHARACTERS FOUND IN ONE TAPE RECORD BY SUBROUTINE
C                   - ALBYTES
C   IHOLD(300)      - THE STRING OF CHARACTERS FOUND BY ALBYTES PACKED FOUR TO A WORD IN
C                   - ZONE DECIMAL FORMAT
C   INTTAB(400)     - TABLE OF INTEGER INTENSITY VALUES RETURNED BY SUBROUTINE TAPERD
C   PNTPER(40)      - TABLE OF THE NUMBER OF INTENSITY READINGS PER PEAK IN THE INTTAB
C                   - LIST
C   DISTAB(40)      - TABLE OF DISTANCE VALUES. ONE VALUE FOR EACH PEAK IN PNTPER
C   PEKNUM          - TOTAL NUMBER OF PEAKS IN THE TAPE RECORD
C   TOTP           - TOTAL NUMBER OF INTENSITY READINGS IN THE TAPE RECORD
C   ERR            - AN ERROR FLAG RETURNED BY SUBROUTINE TAPERD. ERR=0 NORMAL RETURN
C                   - ERR=1 FOR TAPE FORMAT ERROR
C   RAWINT(25000)   - TABLE OF INTENSITY VALUES FOR THE ENTIRE TAPE FILE
C   NUMINT(1000)    - TABLE OF THE NUMBER OF INTENSITY READINGS PER PEAK IN THE ENTIRE
C                   - TAPE FILE
C   RAWDIS(1000)    - TABLE OF DISTANCE READINGS. ONE FOR EACH PEAK IN NUMINT
C   NEXINT          - POINTER TO NEXT VARIABLE IN THE RAWINT ARRAY
C   NEXDIS          - POINTER TO NEXT VARIABLE IN NUMINT AND RAWDIS ARRAYS
C   NOLIST          - LOGICAL VARIABLE TO OPT FOR RAW DATA LISTING. TRUE FOR NO LISTING
C                   - AND FALSE FOR A LISTING
C
C   XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C   LOGICAL NOLIST
C   INTEGER DISTAB,PNTPER,PEKNUM,TOTP,ERR
C   INTEGER RAWINT,NUMINT,RAWDIS
C   COMMON/AMESTP/ IBYTES,IHOLD(300)
C   COMMON/HASPEC/ INTTAB(400),PNTPER(40),DISTAB(40),PEKNUM,TOTP,ERR
C   COMMON/FILE/RAWINT(25000),NUMINT(1000),RAWDIS(1000)
C
C   CLEAR ALL COMMON AREAS
C   DO 500 N=1,25000
C   500  RAWINT(N)=0
C   DO 501 N=1,1000
C   501  NUMINT(N)=0
C   501  RAWDIS(N)=0
C
C   NOLIST IS THE LOGICAL VARIABLE TO OPT FOR A PRINTER LISTING OF THE
C   TAPE FILE
C   READ(5,50) NOLIST
C   NEXINT=1
C   NEXDIS=1
C
C   1000 DO 1001 I=1,300
C   1001  IHOLD(I)=0
C
C   BYTES IS AN AMES LIBRARY SUBROUTINE THAT READS ONE RECORD OF A TAPE
C   C FILE AT A TIME INTO THE IHOLD ARRAY IN ZONE DECIMAL FORMAT. IERR IS
C   AN ERROR FLAG: 0= NORMAL RETURN; 1= END OF FILE; 2=I/O ERROR
C   CALL BYTES(IHOLD,IBYTES,IERR)
C   IF(IERR.EQ.1) GO TO 100
C   IF(IERR.EQ.2) GO TO 1000
C
C   TAPERD IS AN ASSEMBLY SUBROUTINE TO UNBLOCK THE DATA TAPE FORMAT
C   AND CONVERT THE ZONE DECIMAL NUMBERS TO INTEGER. ERR=0 FOR NORMAL
C   RETURN; ERR=1 FOR TAPE FORMAT ERROR
C   1  CALL TAPERD
C   IF(ERR.EQ.1) GO TO 1000
C   2  IF(.NOT.NOLIST) GO TO 300
C   3  IF(PEKNUM.EQ.0) GO TO 4
C   LIMIT1=NEXINT+TOTP-1
C   LIMIT2=NEXDIS+PEKNUM-1
C   KOUNT1=0
C   DO 4001 M=NEXINT,LIMIT1
C   KOUNT1=KOUNT1+1
C   4001  RAWINT(M)=INTTAB(KOUNT1)
C   NEXINT=LIMIT1+1
C   KOUNT2=0
C   DO 4002 M=NEXDIS,LIMIT2
C   KOUNT2=KOUNT2+1
C   NUMINT(M)=PNTPER(KOUNT2)
C   4002  RAWDIS(M)=DISTAB(KOUNT2)
C   NEXDIS=LIMIT2+1
C   GO TO 1000
C
C   UPON READING AN END OF FILE, THE CONTENTS OF THE RAW DATA COMMON
C   BLOCK ARE READ OUT INTO A DISK FILE NAMED IN THE DDEF STATEMENT
C   4  NEXINT=NEXINT-1
C   NEXDIS=NEXDIS-1

```



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C   JANUARY 9,1973                                MICHAEL ROMIEZ
C
      WRITE(7) NEXINT,(RAWINT(I),I=1,NEXINT),NEXDIS,(NUMINT(I),RAWDIS(I),I=1,NEXDIS)
      STOP
C IF AN END OF FILE HAS BEEN READ THE VARIABLE PEKNUM IS SET TO ZERO
100  PEKNUM=0
      GO TO 2
C THIS SECTION WILL PRINT THE CONTENTS OF THE AMES DATA TAPE IF
C REQUESTED BY SETTING NOLIST EQUAL TO FALSE
300  IF(PEKNUM.EQ.0) GO TO 4
      LENGTH=IBYTES/4+1
      INTLN=IBYTES/3
      WRITE(6,16)
      WRITE(6,10) IBYTES
      WRITE(6,11) (IHOLD(J),J=1,LENGTH)
      WRITE(6,17)
      WRITE(6,12) (INTTAB(J),J=1,INTLN)
      WRITE(6,13) PNTPER
      WRITE(6,14) DISTAB
      WRITE(6,15) PEKNUM
      GO TO 3
10  FORMAT(IX,' THE NUMBER OF CHARACTERS IN THIS RECORD ARE ',14)
11  FORMAT(IX,30A4)
12  FORMAT(' ', ' INTENSITIES',30I4)
13  FORMAT(IX,'PNTPER',20I4)
14  FORMAT(IX,'DISTAB',10I8)
15  FORMAT(IX,'PEKNUM',14)
16  FORMAT(IH0)
17  FORMAT(IHX)
50  FORMAT(L5)
      END

```

```

*      JANUARY 9,1973                                MICHAEL ROMIEZ
*
*      THIS SUBROUTINE WILL CONVERT AN AMES HIGH RESOLUTION TAPE RECORD
*      IN EBCDIC FORMAT TO INTEGER FORMAT FOR THE MAIN DATA REDUCTION
*      PROGRAM TO USE.
*
*      XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
TAPE      CSECT
          ENTRY TAPERD
          USING TAPERD,15
TAPERD    SAVE (14,12)
          LR 2,15
          DROP 15
          USING TAPERD,2
* LOAD AMES COMMON BLOCK ADDRESS INTO REG 3
          L 3,COM1
* LOAD MASPEC COMMON BLOCK ADDRESS INTO REG 4
          L 4,COM2
* THIS LOOP WILL CLEAR MASPEC COMMON BLOCK
          LR 5,4
          L 6,=F'482'
          L 7,=F'0'
CLEAR      ST 7,0(5)
          A 5,=F'4'
          BCT 6,CLEAR
* LOAD REGISTER 5 WITH 1BYTES
          L 5,0(0,3)
* ADVANCE AMESTP ADDRESS A FULLWORD TO BEGINNING OF I HOLD ARRAY
          A 3,=F'4'
* INITIALIZE THE TOTAL BYTE COUNTER IN REGISTER 5. THIS IS USED TO TEST
* FOR A NORMAL READ WITH THE FOLLOWING LOGIC. IF WE SUBTRACT SIX FROM
* THE TOTAL BYTES INITALLY AND THREE EVERY TIME WE READ AN INTENSITY
* OR SIX EACH TIME WE READ A DISTANCE THEN WHEN THE COUNTER EQUALS ZERO
* THERE SHOULD ONLY BE SIX ALPHABETIC CHARACTERS LEFT IN THE ARRAY.
* IF THIS IS NOT THE CASE WE TELL THE CALLING PROGRAM THAT THERE IS A
* FORMAT ERROR
          S 5,=F'6'
* LOAD REGISTER SIX AS THE TOTAL OPTICAL DENSITY READING COUNTER
          L 6,=F'0'
* LOAD REGISTER 8 AS THE OPICAL DENSITY PER PEAK COUNTER
          L 8,=F'0'
* LOAD REGISTER 9 WITH THE ADDRESS OF PNTPER
          LR 9,4
          A 9,=F'1600'
* LOAD REG 10 WITH THE ADDRESS OF DISTAB
          LR 10,9
          A 10,=F'160'
* LOAD REG 11 WITH THE ADDRESS OF PEKNUM
          LR 11,10
          A 11,=F'160'
* INITIALIZE REG 12 AS DISTANCE READING COUNTER
          L 12,=F'0'
* WE BEGIN TO PROCESS THE TAPE RECORD
*
* END OF RECORD TEST
*
TESEND    C 5,=F'2'
          BNH EXIT
* WE NOW ENTER THE TAPE READ LOGIC
*
TESTCH    TM 0(3),X'F0'
          BNO DISTCE
          TM 2(3),X'F0'
          BNO ERROR
* IF THE CHARACTER IS NUMERIC IT IS AN INTENSITY READING
* WE CONVERT IT TO BINARY AND STORE IT IN INTTAB
          PACK DBL,0(3,3)
* IF THE INTENSITY IS NEGITIVE WE HAVE A TAPE FORMAT ERROR
          CP DBL,=P'0'
          BNH ERROR
          CVB 7,DBL
          ST 7,0(0,4)
* INCREMENT THE COUNTERS AND DATA BASE REGISTERS
          A 3,=F'3'
          A 4,=F'4'
          A 8,=F'1'
          A 6,=F'1'
* SUBTRACT THREE FROM TOTAL BYTE COUNTER
          S 5,=F'3'
          B TESEND
* IF THE CHARACTER IS ALPHABETIC IT IS A DISTANCE READING
* WE CONVERT IT TO BINARY AND STORE IT IN DISTAB
DISTCE    TM 5(3),X'D0'

```

```

*   JANUARY 9,1973                               MICHAEL ROMIEZ
*
      BNO   ERROR
      PACK DBL,0(6,3)
* CONVERT THE DISTANCE READING TO A LEGITIMATE POSITIVE NUMBER
      OI    DBL+7,X'0F'
      CVB   7,DBL
      ST    7,0(0,10)
* ADVANCE THE DISTAB BASE REGISTER AND ADD 1 TO THE DISTANCE READING
* COUNTER,SUBTRACT SIX FROM THE TOTAL BYTE COUNTER
      A     10,=F'4'
      A     12,=F'11'
      A     3,=F'6'
      S     5,=F'6'
* STORE THE NUMBER OF OPTICAL DENSITY READINGS FOR THIS PEAK AND
* ADVANCE THE PNTPER BASE REGISTER
      ST    8,0(0,9)
      A     9,=F'4'
* RE INITIALIZE THE INTENSITY NUMBER COUNTER
      L     8,=F'0'
      B     TESEND
* IF TAPE FORMAT IS INCORRECT GO TO ERROR EXIT
*
EXIT   TM    0(3),X'F0'
      BO    ERROR
      TM    5(3),X'F0'
      BO    ERROR
* FOR CORRECT TAPE FORMAT TAKE NORMAL EXIT
*
* STORE FINAL DISTANCE READING IN DISTAB
      PACK DBL,0(6,3)
      OI    DBL+7,X'0F'
      CVB   7,DBL
      ST    7,0(0,10)
* INCREMENT DISTANCE READING COUNTER
      A     12,=F'11'
* STORE THE NUMBER OF OPTICAL DENSITY READINGS IN THE LAST RECORD
* IN PNTPER
      ST    8,0(0,9)
* STORE THE NUMBER OF PEAKS IN THIS RECORD IN PEKNUM
      ST    12,0(0,11)
* WE ADD FOUR TO THE ADDRESS OF PEKNUM TO GET TO TOTP
      A     11,=F'4'
* STORE THE TOTAL NUMBER OF OPTICAL DENSITY READINGS IN THIS RECORD
* IN TOTP
      ST    6,0(0,11)
* WE ADD FOUR TO THE ADDRESS OF TOTP TO GET TO THE ERROR FLAG
      A     11,=F'4'
* THIS IS A NORMAL EXIT STORE ZERO IN ERROR FLAG
      L     4,=F'0'
      ST    4,0(0,11)
      B     OUT
* WE ADD EIGHT TO THE ADDRESS OF PEKNUM TO GET TO THE ERROR FLAG
ERROR  A     11,=F'8'
* THIS IS AN ABNORMAL EXIT STORE ONE IN THE ERROR FLAG
      L     4,-F'11'
      ST    4,0(0,11)
      B     OUT
* RETURN CONTROL TO CALLING PROGRAM
OUT    RETURN (14,12)
      BR    14
DBL    DS    D
COM1   DC    V(AMESTP)
COM2   DC    V(MASPEC)
      END    TAPE

```

```
C      OCTOBER 10,1972                      MICHAEL ROMIEZ
C
C      THIS IS THE PROGRAM WHICH CALLS THE VARIOUS SUBROUTINES TO REDUCE THE HIGH
C      RESOLUTION MASS SPECTRAL DATA.
C
C      XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C      CALL PARLST
C      CALL PKLIST
C      CALL STDFIT
C      CALL EELST
C      STOP
C      END
```



```

JANUARY 30,1973                                MICHAEL ROMEZ
C THIS SUBROUTINE IS A GENERAL POLYNOMIAL FITTING ALGORITHM. IT WILL FIT UP TO 200
C DATA POINTS TO A FROM FIRST TO TENTH DEGREE POLYNOMIAL. THE POLYNOMIAL IS BEST FIT
C IN THE LEAST SQUARES SENSE.
C
C VARIABLES
C X(200) - UP TO 200 ABSCISSA VALUES
C Y(200) - UP TO 200 ORDINATE VALUES
C C(11) - UP TO 11 COEFFICIENTS CALCULATED BY THIS SUBROUTINE
C N - THE NUMBER OF DATA POINTS IN THE X AND Y ARRAYS
C IDEG - THE DEGREE OF THE POLYNOMIAL
C SUMX(20) - THE UP TO TWENTY VALUES OF THE SUMMATIONS OF X RAISED TO THE
C POWERS OF ONE THROUGH TWICE IDEG INCLUSIVE.
C SUMXY(10) - THE UP TO TEN VALUES OF THE SUMMATIONS OF Y TIMES X RAISED
C TO THE POWERS OF ONE THROUGH IDEG INCLUSIVE.
C SUMY - THE SUMMATION OF THE Y VALUES
C A(11,11) - THE IDEG+1 BY IDEG+1 SQUARE MATRIX OF SUMMATION VALUES FROM
C THE NORMAL EQUATIONS OF THE LEAST SQUARE SYSTEM OF EQUATIONS
C B(11) - THE UP TO ELEVEN CONSTANT TERMS OF THE NORMAL EQUATIONS
C DM(11) - TEMP STORAGE
C
C XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
SUBROUTINE POLYFT
DOUBLE PRECISION B(11),A(11,11),C(11),DM(11)
DOUBLE PRECISION SUMX(20),SUMXY(10),SUMY,X(200),Y(200)
COMMON/FIT/X,Y,N,IDEG,C
C CLEAR VARIABLES
DO 1000 J=1,10
1000 SUMXY(J)=0.
DO 1001 J=1,11
B(J)=0.
DM(J)=0.
1001 C(J)=0.
DO 1002 J=1,20
1002 SUMX(J)=0.
DO 1003 K=1,11
DO 1003 K=1,11
1003 A(J,K)=0.
SUMY=0.
C CALCULATE THE SUMMATION VALUES
DO 2000 M=1,IDEG
DO 2000 K=1,N
2000 SUMXY(M)=SUMXY(M)+Y(K)*X(K)**M
DO 2001 M=1,N
2001 SUMY=SUMY+Y(M)
N2=IDEG**2
DO 2002 M=1,N2
DO 2002 K=1,N
2002 SUMX(M)=SUMX(M)+X(K)**M
C THIS SECTION ASSIGNS THE SUMMATION VALUES TO THE PROPER
C MATRIX ELEMENTS
LIMIT=IDEG+1
DO 3001 J=1,LIMIT
DO 3000 K=1,LIMIT
M=J+K-2
IF (M.EQ.0) GO TO 3000
A(J,K)=SUMX(M)
3000 CONTINUE
IF (J.EQ.1) GO TO 3001
B(J)=SUMXY(J-1)
3001 CONTINUE
A(1,1)=N
B(1)=SUMY
C THIS SECTION CALCULATES THE ELEMENTS OF THE TRIANGULAR MATRIX
NN=0
DO 4001 J=1,IDEG
NN=NN+1
DO 4001 K=NN,IDEG
DM(K)=A(K+1,J)/A(J,J)
DO 4000 L=1,LIMIT
4000 A(K+1,L)=A(K+1,L)-DM(K)*A(J,L)
B(K+1)=B(K+1)-DM(K)*B(J)
4001 CONTINUE
C THIS SECTION CALCULATES THE COEFFICIENTS OF THE CURVE FITTING
C POLYNOMIAL FROM THE TRIANGULAR MATRIX
M=LIMIT+1
DO 5001 K=1,LIMIT
DM(1)=0.D0
IROW=M-K
ICOL=M-K
5000 ICOL=ICOL+1

```

C JANUARY 30, 1973

MICHAEL ROMIEZ

C
C IF(ICOL.GT.LIMIT) GO TO 5001
DM(1)=DM(1)-A(IROW,ICOL)*C(ICOL)
GO TO 5000
5001 C(IROW)=(B(IROW)+DM(1))/A(IROW,IROW)
RETURN
END

C FEBRUARY 5,1973

MICHAEL ROMIEZ

C THIS MODULE SETS ALL THE VARIABLE PARAMETERS USED BY THE DECONVOLUTION, PEAK CENTER
C DETERMINATION, AND DISTANCE TO MASS CONVERSION ROUTINES.

C THE VARIABLES HAVE DEFALT VALUES GIVEN AT COMPILATION, BUT THEY MAY BE REVIEWED AND
C CHANGED AT EXECUTION TIME.

C VARIABLES

C DERDEG - THE DEGREE OF THE FIRST DERIVATIVE POLYNOMIAL
C DERNUM - THE NUMBER OF DATA POINTS USED BY THE FIRST DERIVATIVE POLYNOMIAL
C SMNUM - THE NUMBER OF DATA POINTS USED BY THE SMOOTHING POLYNOMIAL
C SMDEG - THE DEGREE OF THE SMOOTHING POLYNOMIAL
C CONNUM - THE NUMBER OF POINTS USED BY THE DISTANCE TO MASS CONVERSION POLYNOMIAL
C CONDEG - THE DEGREE OF THE DISTANCE OF MASS CONVERSION POLYNOMIAL
C CUTOFF - IF THE TOTAL NUMBER OF DATA POINTS IN THE PEAK PROFILE IS LESS THAN OR
C EQUAL THIS NUMBER, THE PEAK IS TREATED AS A SINGLE PEAK WITH NO
C DECONVOLUTION ANALYSIS
C INTMIN - IF THE PEAK PROFILE CONTAINS NO INTENSITY READINGS ABOVE THIS VALUE, THE
C PEAK IS IGNORED. THIS ABILITY IS USEFUL IN DISCRIMINATION AGAINST SUCH
C ARTIFACTS AS SCRATCHES ON THE PHOTOPLATE
C TOLFAC - THIS IS A FACTOR USED BY THE STANDARD MASS SEARCHING ROUTINE TO CALCULATE
C THE TOLERANCE WITHIN WHICH IT WILL ACCEPT A PEAK AS A STANDARD MASS.
C EXPTOL - LOWER LIMIT OF THE NUMBER OF INTENSITY VALUES CONSIDERED AN
C OVEREXPOSURE.

C XXX

SUBROUTINE PARLST
INTEGER DERDEG,DERNUM,SMNUM,SMDEG,CUTOFF,CONNUM,CONDEG,TOLFAC,EXPTOL
INTEGER YES/'Y'/
COMMON/PAR/DERDEG,DERNUM,SMDEG,SMNUM,CUTOFF,INTMIN,CONNUM,CONDEG,TOLFAC,EXPTOL
DERDEG=3
DERNUM=5
SMNUM=25
SMDEG=3
CONNUM=4
CONDEG=2
CUTOFF=30
INTMIN=90
TOLFAC=7
EXPTOL=180
WRITE(6,100)
100 FORMAT(1X'DO YOU WANT TO REVIEW THE PEAK POSITION PARAMETERS?')
READ(6,101) I
101 FORMAT(A1)
IF(I.NE.YES) RETURN
WRITE(6,102) DERDEG
102 FORMAT(1X,'THE DEGREE OF THE FIRST DERIVATIVE POLYNOMIAL IS',13)
CALL INTGER(M)
IF(M.NE.0) DERDEG=M
WRITE(6,103) DERNUM
103 FORMAT(1X,'THE NUMBER OF POINTS USED FOR THE FIRST DERIVATIVE IS',13)
CALL INTGER(M)
IF(M.NE.0) DERNUM=M
WRITE(6,104) SMNUM
104 FORMAT(1X,'THE NUMBER OF POINTS USED FOR THE SMOOTHING POLYNOMIAL IS',13)
CALL INTGER(M)
IF(M.NE.0) SMNUM=M
WRITE(6,105) SMDEG
105 FORMAT(1X,'THE DEGREE OF THE POLYNOMIAL USED FOR SMOOTHING IS',13)
CALL INTGER(M)
IF(M.NE.0) SMDEG=M
WRITE(6,108) CONNUM
108 FORMAT(1X,'THE NUMBER OF MASSES USED BY THE MASS CALCULATION POLYNOMIAL IS',13)
CALL INTGER(M)
IF(M.NE.0) CONNUM=M
WRITE(6,109) CONDEG
109 FORMAT(1X,'THE DEGREE OF THE MASS CALCULATION POLYNOMIAL IS',13)
CALL INTGER(M)
IF(M.NE.0) CONDEG=M
WRITE(6,106) CUTOFF
106 FORMAT(1X,'THE CUTOFF VALUE FOR A SINGLE PEAK IS',14)
CALL INTGER(M)
IF(M.NE.0) CUTOFF=M
WRITE(6,107) INTMIN
107 FORMAT(1X,'THE MINIMUM HIGH INTENSITY FOR A REAL PEAK IS',14)
CALL INTGER(M)
IF(M.NE.0) INTMIN=M
WRITE(6,110) TOLFAC
110 FORMAT(1X,'THE SLIDING TOLERANCE FACTOR FOR STANDARD MASS IDENTIFICATION IS',13)
CALL INTGER(M)
IF(M.NE.0) TOLFAC=M

C FEBRUARY 5, 1973
C

MICHAEL ROMIEZ

```
WRITE(6,111) EXPTOL
111 FORMAT(1X,'THE NUMBER OF INTENSITY VALUES ABOVE WHICH A PEAK WILL BE EDITED IS',I4)
CALL INTGER(M)
IF(M.NE.0) EXPTOL=M
RETURN
END
```



```

JANUARY 11,1973                                MICHAEL ROMIEZ
THIS SUBROUTINE SENDS RAW PEAK PROFILES TO SUBROUTINE PEAK WHICH IN TURN CONTROLS
THE SMOOTHING, DECONVOLUTION, AND PEAK CENTER CALCULATION ROUTINES. THIS MODULE
SENDS UP TO 400 DATA POINTS TO SUBROUTINE PEAK AND IS RETURNED UP TO TEN PEAK CENTERS
AND MAXIMUM INTENSITY VALUES. THESE CENTERS AND INTENSITIES ARE STORED FOR THE REST
OF THE DATA REDUCTION ROUTINES.

VARIABLES
RAWINT(25000)  - TABLE OF INTENSITY VALUES FOR THE ENTIRE RAW DATA FILE
NUMINT(1000)   - TABLE OF THE NUMBER OF INTENSITY READINGS PER PEAK IN THE ENTIRE
                - RAW DATA FILE
RAWDIS(1000)   - TABLE OF DISTANCE READINGS. ONE FOR EACH PEAK IN NUMINT
CENTER(1500)   - DISTANCE OF PEAK CENTERS FROM THE LEFT EDGE OF THE PHOTOPLATE
                - IN MILLIMETERS
HIGHT(1500)    - MAXIMUM INTENSITY VALUE OF EACH PEAK
NUMPK          - NUMBER OF PEAKS IN THE PEAK LIST TABLE
LIST(400)      - HOLDING TABLE TO PASS ONE PEAK PROFILE BETWEEN MODULES
NUM            - EITHER THE NUMBER OF INTENSITY VALUES IN THE PEAK PROFILE OR THE
                - NUMBER OF PEAK CENTERS RETURNED BY SUBROUTINE PEAK
DIS           - THE DISTANCE IN HALF-MICRONS OF THE LAST INTENSITY READING IN THE
                - PEAK PROFILE MEASURED FROM THE LEFT END OF THE PHOTOPLATE
CENT(10)       - UP TO 10 PEAK CENTERS RETURNED TO THIS SUBROUTINE FOR EACH PEAK
                - PROFILE
HGT(10)        - UP TO TEN MAXIMUM INTENSITY VALUES, ONE FOR EACH PEAK IN CENT
LSTPAR(10)     - LIST OF PARAMETERS FROM THE PARLIST MODULE
PTINT          - POINTS TO NEXT INTENSITY READING IN RAWINT
PTDIS         - POINTS TO NEXT POSITION IN THE RAWDIS AND NUMINT ARRAYS
PTCENT        - POINTS TO NEXT POSITION IN THE CENTER AND HIGHT ARRAYS
INTMIN        - ALL INTENSITY READINGS IN A PROFILE MUST BE ABOVE THIS VALUE
                - OR THE PEAK IS IGNORED. THIS IS USEFUL IN DISCRIMINATING AGAINST
                - SCRATCHES ON THE PHOTOPLATE
IEDIT         - LOWER LIMIT OF THE NUMBER OF INTENSITY READINGS FOR EDITING

XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
SUBROUTINE PKLIST
DOUBLE PRECISION CENTER,CENT(10)
INTEGER RAWINT(25000),NUMINT(1000),RAWDIS(1000)
INTEGER PTDIS,PTINT,LIST(400),PTCENT,DIS
INTEGER YES/'Y'/
DIMENSION HGT(10)
COMMON/PKLIST/ CENTER(1500),HIGHT(1500),NUMPK
COMMON/PK/ LIST,NUM,DIS,CENT,HGT
COMMON/PAR/ LSTPAR(10)
C SET THE INTMIN VALUE AS GIVEN BY THE PARAMETER LIST MODULE
    INTMIN=LSTPAR(6)
C SET THE LOWER LIMIT OF THE NUMBER OF INTENSITY READINGS FOR PROFILE EDITING
    IEDIT=LSTPAR(10)
C CLEAR VARIABLES
    DO 100 M=1,1500
        CENTER(M)=0.
100  HIGHT(M)=0.
    DO 150 M=1,25000
150  RAWINT(M)=0
    DO 151 M=1,1000
        RAWDIS(M)=0
151  NUMINT(M)=0
C READ IN RAW DATA FROM DISK FILE
    READ(9) N,(RAWINT(I),I=1,N),M,(NUMINT(I),RAWDIS(I),I=1,M)
    PTINT=0
    PTDIS=1
    PTCENT=0
    DO 10 I=1,10
10  HGT(I)=0.
    DO 15 I=1,10
15  CENT(I)=0.
C RETURN IF THERE ARE NO MORE PROFILES IN THE RAW DATA FILE
    IF(NUMINT(PTDIS).EQ.0) GO TO 9999
C IF THERE ARE MORE THAN 400 POINTS IN A PARTICULAR PEAK PROFILE, THE PEAK IS
C IGNORED. A PEAK 200 MICRONS WIDE IS TOO WIDE TO BE A REAL PEAK.
    IF(NUMINT(PTDIS).GT.400) GO TO 11
    IF(NUMINT(PTDIS).GT.4) GO TO 200
C IF A PROFILE HAS FEWER THAN 5 INTENSITY READINGS IT IS DELETED AS A NOISE SPIKE
11  PTDIS=PTDIS+1
    PTINT=PTINT+NUMINT(PTDIS-1)
    GO TO 10
C READ PEAK PROFILE AND DISTANCE INTO DATA PASSING ARRAYS
200 NUM=NUMINT(PTDIS)
    DO 225 I=1,400
225 LIST(I)=0
    DO 250 J=1,NUM
        PTINT=PTINT+1
250 LIST(J)=RAWINT(PTINT)
        DIS=RAWDIS(PTDIS)

```

```

C   JANUARY 11,1973                                MICHAEL ROMIEZ
C
C IF THE HIGHEST INTENSITY VALUE IS LESS THAN INTMIN IGNORE THE PEAK
DO 275 J=1,NUM
  IF(LIST(J).GT.INTMIN) GO TO 290
275 CONTINUE
  PTDIS=PTDIS+1
  GO TO 10
C IF THERE ARE MORE THAN IEDIT INTENSITY READINGS IN A PEAK PROFILE, IT IS AN
C OVEREXPOSURE AND IS DISPLAYED FOR POSSIBLE EDITING.
290 IF(NUM.GT.IEDIT) CALL EDIT
C SEND PEAK PROFILE TO BE ANALYZED
  CALL PEAK
C KEEP PEAK CENTERS AND INTENSITIES IN CENTER AND HIGHT ARRAYS
DO 299 M=1,NUM
  PTCENT=PTCENT+1
C THE PEAK CENTER VALUES ARE CONVERTED FROM HALF-MICRONS TO MILLIMETERS
  CENTER(PTCENT)=CENT(M)/2000.
C THE DISTANCE ENCODER RECYCLES AT 255 MILLIMETERS AND THIS TEST
C COMPENSATES FOR RECYCLING.
  IF(PTCENT.EQ.1) GO TO 299
  IF(CENTER(PTCENT).LT.CENTER(PTCENT-1).AND.CENTER(PTCENT-1)-
    X.GT.200) CENTER(PTCENT)=CENTER(PTCENT)+255.
299 HEIGHT(PTCENT)=HGT(M)
C THIS TEST WILL TRUNCATE THE PEAK LIST IF IT APPROACHES WITHIN 10 VARIABLES OF
C THE END OF THE ARRAY
  IF(PTCENT.GT.1490) GO TO 400
  PTDIS=PTDIS+1
  GO TO 10
400 WRITE(6,1)
1  FORMAT('0','THE PEAK LIST WAS TRUNCATED AT',16,' VALUES')
9999 NUPK=PTCENT
  RETURN
  END

```

```

C   JANUARY 17,1973                               MICHAEL ROMIEZ
C
C   THIS SUBROUTINE CONTROLS DATA SMOOTHING, PEAK DECONVOLUTION, AND PEAK CENTER CALCULATION
C
C                                     VARIABLES
C
C   LIST(400)      - HOLDING TABLE TO PASS ONE PEAK PROFILE BETWEEN MODULES
C   NUM            - EITHER THE NUMBER OF INTENSITY VALUES IN THE PEAK PROFILE OR THE
C                  - NUMBER OF PEAK CENTERS RETURNED BY SUBROUTINE ANAL
C   DIS            - THE DISTANCE IN HALF-MICRONS OF THE LAST INTENSITY READING IN THE
C                  - PEAK PROFILE MEASURED FROM THE LEFT END OF THE PHOTOPLATE
C   CENT(10)       - UP TO 10 PEAK CENTERS RETURNED TO THIS SUBROUTINE FOR EACH PEAK
C                  - PROFILE
C   HGT(10)        - UP TO TEN MAXIMUM INTENSITY VALUES, ONE FOR EACH PEAK IN CENT
C   LSTPAR(10)     - LIST OF PARAMETERS FROM THE PARLIST MODULE
C   NUMSM          - NUMBER OF POINTS USED BY THE SMOOTHING ROUTINE
C   CUTOFF         - MINIMUM NUMBER OF POINTS FOR A PROFILE TO BE ANALYZED FOR
C                  - MORE THAN ONE PEAK. PROFILES WITH LESS THAN CUTOFF DATA
C                  - POINTS WILL BE TREATED AS A SINGLE PEAK
C
C   XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C   SUBROUTINE PEAK
C   DOUBLE PRECISION CENT(10)
C   INTEGER DIS,CUTOFF
C   DIMENSION HGT(10),LIST(400)
C   COMMON/PAR/LSTPAR(10)
C   COMMON/PK/LIST,NUM,DIS,CENT,HGT
C   SET THE PARAMETERS FOR THE NUMBER OF POINTS USED BY THE SMOOTHING ROUTINE AND
C   THE MINIMUM NUMBER OF POINTS IN A PROFILE CONSIDERED FOR DECONVOLUTION ANALYSIS
C   NUMSM=LSTPAR(4)
C   CUTOFF=LSTPAR(5)
C   IF THERE ARE LESS THAN CUTOFF DATA POINTS, THIS IS A SINGLE PEAK
C   IF(NUM.GT.CUTOFF) GO TO 100
C   IF THERE ARE ENOUGH RAW DATA POINTS IN THE PROFILE, SMOOTH THE DATA
C   BEFORE SENDING IT TO MODULE ONEPK
C   IF(NUM.GT.(NUMSM+10)) GO TO 150
C   CALL ONEPK
C   RETURN
C   150 CALL SMOOTH
C   THE DATA SMOOTHING ROUTINE RETURNS (NUMSM-1) LESS DATA POINTS THAN WAS IN THE
C   RAW DATA. THE DISTANCE VALUE WHICH INDICATES THE DISTANCE OF THE LAST DATA POINT
C   IN THE LIST MUST BE ADJUSTED
C   DIS=DIS-((NUMSM-1)/2)
C   CALL ONEPK
C   RETURN
C   SMOOTH THE DATA
C   100 CALL SMOOTH
C   THE DATA SMOOTHING ROUTINE RETURNS (NUMSM-1) LESS DATA POINTS THAN WAS IN THE
C   RAW DATA. THE DISTANCE VALUE WHICH INDICATES THE DISTANCE OF THE LAST DATA POINT
C   IN THE LIST MUST BE ADJUSTED
C   DIS=DIS-((NUMSM-1)/2)
C   SUBROUTINE ANAL USES THE FIRST DERIVATIVE TO DETERMINE IF THERE IS MORE THAN ONE
C   PEAK IN A PARTICULAR SMOOTHED PROFILE
C   CALL ANAL
C   RETURN
C   END

```



```

C   JANUARY 16,1973                                MICHAEL ROMIEZ
C
      J=1
      M=0
      1 M=M+1
      IF (M.GT.INDEX3-1) GO TO 99
C UPSLOPE LOGIC
      IF (FSTDER(M+1)-FSTDER(M))3,1,1
      2 M=M+1
      IF (M.GT.INDEX3) GO TO 99
C WE HAVE JUST STARTED DOWN THE SIDE OF THE FIRST DERIVATIVE CURVE. WE CONTINUE DOWN
C UNTIL WE GO THROUGH THE X-AXIS. THE POINT AT WHICH THE FIRST DERIVATIVE CURVE PASSES
C THROUGH THE X-AXIS GOING FROM POSITIVE TO NEGATIVE, IS THE CENTER OF A PEAK.
      3 IF (FSTDER(M).GT.0.) GO TO 2
C THE NEXT SIX STATEMENTS CALCULATE THE CENTER AND HEIGHT OF THE JUST FOUND PEAK
      DHIGH=FSTDER(M-1)
      DLOW=ABS(FSTDER(M))
      CENT(J)=FLOAT(M-1)+DHIGH/DHIGH+DLOW
      CENT(J)=CENT(J)+LIMIT/2
      K=M+LIMIT/2
      HGT(J)=LIST(K)
C IF THIS MODULE HAS ALREADY FOUND TEN PEAKS, STOP LOOKING
      J=J+1
      IF (J.GT.10) GO TO 99
C THIS LOOP TESTS IF THE FIRST DERIVATIVE CURVE GOES BACK ABOVE THE X-AXIS. IF IT
C DOES IT CAUSES A BRANCH TO THE UPSLOPE LOGIC SECTION
      DO 80 K=M,INDEX3
      IF (FSTDER(K).LT.0.) GO TO 80
      GO TO 101
      80 CONTINUE
      GO TO 99
      101 M=K-1
      GO TO 1
      99 NN=J-1
C THE CENTER OF EACH PEAK FOUND BY THIS MODULE IS CALCULATED IN TERMS OF THE DISTANCE
C IN HALF-MICRONS FROM THE LEFT EDGE OF THE PHOTO PLATE.
      DO 89 K=1,NN
      89 CENT(K)=DIS-(NUM-CENT(K))
C NUM IS SET TO THE NUMBER OF PEAKS FOUND
      NUM=NN
      RETURN
      END

```



```

C      JANUARY 17,1973                                MICHAEL ROMIEZ
C
C      THIS SUBROUTINE WILL DETERMINE THE CENTER AND HIGHT OF A SINGLE PEAK. IT FITS
C      THE PEAK PROFILE TO A GAUSSIAN DISTRIBUTION CURVE.
C
C
C      VARIABLES
C
C      X(200)      - UP TO 200 POINTS THAT ARE USED BY THE POLYNOMIAL FITTING
C                  - ROUTINE FOR THE BEST LEAST SQUARES GAUSSIAN FIT
C      Y(200)      - ONE ORDINATE VALUE FOR EACH X VALUE
C      N           - NUMBER OF X AND Y VALUES
C      IDEG        - IN THIS CASE IDEG=2. IT IS THE DEGREE OF THE POLYNOMIAL USED
C                  - FOR FITTING
C      C(11)       - THE FIRST THREE VALUES OF THE ARRAY CONTAIN THE COEFFICIENTS OF
C                  - THE SECOND DEGREE POLYNOMIAL
C      LIST(400)   - HOLDING TABLE TO PASS PEAK PROFILES BETWEEN MODULES
C      NUM         - EITHER THE NUMBER OF INTENSITIES IN THE PEAK PROFILE OR NUM=1
C                  - TO INDICATE THE NUMBER OF PEAKS FOUND BY THIS MODULE
C      DIS         - THE DISTANCE IN HALF-MICRONS OF THE LAST INTENSITY READING IN
C                  - THE PROFILE AS MEASURED FROM THE LEFT END OF THE PHOTOGRAPHIC
C                  - PLATE
C      CENT(10)    - THE CENTER CALCULATED BY THIS MODULE IS PUT IN CENT(1)
C      HGT(10)     - THE HEIGHT CALCULATED BY THIS MODULE IS PUT IN HGT(1)
C
C      XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C      SUBROUTINE ONEPK
C      DOUBLE PRECISION CENT(10),C(11),X(200),Y(200)
C      INTEGER DIS
C      DIMENSION HGT(10),LIST(400)
C      COMMON/FIT/X,Y,N,IDEG,C
C      COMMON/PK/LIST,NUM,DIS,CENT,HGT
C      IF(NUM.GT.200) RETURN
C      N=NUM
C      IDEG=2
C
C      PUT THE LOG OF THE INTENSITY VALUE IN THE ORDINATE LIST
C      DO 50 J=1,NUM
C      X(J)=DFLOAT(J)
C      50 Y(J)=DLOG(DFLOAT(LIST(J)))
C
C      CALL THE POLYNOMIAL FITTING ROUTINE
C      CALL POLYFT
C      IF(C(3).GT.0.) GO TO 99
C
C      CALCULATE THE CENTER AND HEIGHT OF THE PEAK FROM THE COEFFICIENTS OF THE POLYNOMIAL
C      CENT(1)=(-C(2)/C(3))/2.
C      HGT(1)=C(1)+(CENT(1)/SQRT(-1./C(3)))**2
C      HGT(1)=EXP(HGT(1))
C      CENT(1)=DIS-(NUM-CENT(1))
C      NUM=1
C      99 RETURN
C      END

```

```

C      FEBRUARY 3,1973                                MICHAEL ROMIEZ
C
C      THIS MODULE WILL ALLOW EDITING OF THE RAW PEAK PROFILE. PART OF THE PROFILE MAY BE
C      DELETED AND SELECTED INTENSITY READINGS MAY BE CHANGED.
C
C
C      VARIABLES
C
C      LIST(400)      - HOLDING TABLE TO PASS ONE PEAK PROFILE BETWEEN MODULES
C      NUM            - THE NUMBER OF INTENSITY VALUES IN THE PEAK PROFILE
C      DIS            - THE DISTANCE IN HALF-MICRONS OF THE LAST INTENSITY READING IN THE
C                      - PEAK PROFILE MEASURED FROM THE LEFT END OF THE PHOTOPLATE
C      CENT(10)       - UP TO 10 PEAK CENTERS
C      HGT(10)        - UP TO TEN MAXIMUM INTENSITY VALUES, ONE FOR EACH PEAK IN CENT
C
C      XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C      SUBROUTINE EDIT
C      DOUBLE PRECISION CENT(10)
C      INTEGER LIST(400),DIS
C      DIMENSION HGT(10)
C      COMMON/PK/LIST,NUM,DIS,CENT,HGT
C      DATA IYES/'Y'/
C
C      THIS SECTION LISTS THE RAW PEAK PROFILE
C      WRITE(6,100)
C      100 FORMAT('0','PEAK PROFILE')
C      WRITE(6,101)(LIST(I),I=1,NUM)
C      101 FORMAT(5X,2015)
C
C      THIS SECTION WILL DELETE PART OF THE PEAK PROFILE. IT WILL ACCEPT A LOW AND HIGH
C      SUBSCRIPT AND ADJUST THE INTENSITY READINGS IN ARRAY LIST TO THESE NEW VALUES.
C      IF THE NEW LOW SUBSCRIPT ENTERED IS GREATER THAN NUM, THIS SECTION WILL BE SKIPPED.
C      WRITE(6,102)
C      102 FORMAT('0','ENTER A NEW STARTING INTENSITY')
C      CALL INTGER(LOW)
C      IF(LOW.GT.NUM) GO TO 400
C      IF (I.EQ.1) GO TO 200
C      LIMIT=NUM-(LOW-1)
C      DO 150 J=1,LIMIT
C      INDEX=LOW-1+J
C      150 LIST(J)=LIST(INDEX)
C      200 WRITE(6,104)
C      104 FORMAT('0','ENTER NEW ENDING SUBSCRIPT')
C
C      IF NOTHING IS ENTERED FOR THE ENDING SUBSCRIPT, THE OLD VALUE IS KEPT.
C      CALL INTGER(IHIGH)
C      IF(IHIGH.EQ.0) IHIGH=NUM
C      IF(IHIGH.EQ.NUM) GO TO 300
C      ISTRT=IHIGH+1
C      DO 250 J=ISTRT,NUM
C      250 LIST(J)=0
C      DIST=DIS-(NUM-IHIGH)
C      300 NUM=NUM-(NUM-IHIGH)-(LOW-1)
C
C      THIS SECTION WILL ACCEPT INTENSITY VALUE CHANGES
C      400 WRITE(6,105)
C      105 FORMAT('0','ANY INTENSITY EDITING?')
C      READ(5,106) I
C      106 FORMAT(A1)
C      IF(I.NE.IYES) RETURN
C      WRITE(6,101)(LIST(I),I=1,NUM)
C      405 WRITE(6,107)
C      107 FORMAT(IX,'ENTER SUBSCRIPT')
C      CALL INTGER(M)
C      IF(M.EQ.0) RETURN
C      WRITE(6,108)
C      108 FORMAT(IX,'ENTER INTENSITY')
C      CALL INTGER(N)
C      LIST(M)=N
C      GO TO 450
C      END

```

```

C      JANUARY 26,1973                                MICHAEL ROMIEZ
C
C      THIS SUBROUTINE CONTROLS THE STANDARD MASS SEARCH AND LISTING ROUTINES
C
C
C
C      VARIABLES
C
C      CENTER(1500)  - DISTANCE OF PEAK CENTERS FROM THE LEFT EDGE OF THE PHOTOPLATE
C                   - IN MILLIMETERS
C      HIGHT(1500)   - MAXIMUM INTENSITY VALUE OF EACH PEAK
C      NUMPK         - NUMBER OF PEAKS IN THE PEAK LIST TABLE
C      STDDIS(160)   - THE DISTANCE VALUES OF PEAK CENTERS IDENTIFIED AS STANDARD MASSES
C      STDMA(160)    - THE SQUARE ROOT OF THE MASS VALUE OF IDENTIFIED STANDARD MASSES
C      NUMSTD        - THE NUMBER OF IDENTIFIED STANDARD MASSES
C      REFMA(160)    - A TABLE OF THE MOST INTENSE LINES IN THE MARKER COMPOUND SPECTRUM.
C                   - THESE ARE THE MASSES SEARCHED FOR BY THE STANDARD MASS IDENTIFICATION
C                   - ROUTINE
C
C      XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C      SUBROUTINE STDFIT
C      DOUBLE PRECISION CENTER,STDDIS(160),STDMA(160),REFMA(160)
C      INTEGER YES/'Y'/
C      COMMON/PKLST/ CENTER(1500),HIGHT(1500),NUMPK
C      COMMON/PFKLST/ STDDIS,STDMA,REFMA,NUMSTD
C
C      CLEAR VARIABLES
C      DO 99 M=1,160
C      STDDIS(M)=0.
C      STDMA(M)=0.
C      99 STDMA(M)=0.
C      READ IN THE STANDARD MASSES AND GET THEIR SQUARE ROOT
C      READ(7) REFMA
C      DO 100 M=1,160
C      100 REFMA(M)=DSQRT(REFMA(M))
C      LIST UP TO THE FIRST FORTY DISTANCE VALUES IN ARRAY CENTER ON THE OUTPUT TERMINAL
C      I=NUMPK
C      IF(I.GT.40) I=40
C      WRITE(6,1) I
C      1 FORMAT('0','THESE ARE THE FIRST',I3,'DISTANCE VALUES')
C      WRITE(6,2) (CENTER(J),J=1,I)
C      2 FORMAT(1X,10F12.5)
C      NUMSTD=0
C      3 NUMSTD=NUMSTD+1
C      C THE PROGRAM USER MUST ENTER A KNOWN MASS VALUE FOR AT LEAST TWO OF THE MASS VALUES
C      C IN ARRAY CENTER.
C      WRITE(6,4)
C      4 FORMAT(1X,'ENTER SUBSCRIPT')
C      CALL INTGER(I)
C      IF(I.EQ.0) GO TO 66
C      WRITE(6,5)
C      5 FORMAT(1X,'ENTER MASS')
C      READ(5,6) STDMA(NUMSTD)
C      6 FORMAT(F10.0)
C      STDDIS(NUMSTD)=CENTER(I)
C      STDMA(NUMSTD)=DSQRT(STDMA(NUMSTD))
C      GO TO 3
C      66 NUMSTD=NUMSTD-1
C      C AT THIS POINT THE USER CAN TRANSFER CONTROL TO THE STANDARD MASS MANUAL ADDITION MODULE
C      C SO THAT HE CAN USE THE LIST EDITING CAPABILITIES OF THAT MODULE.
C      WRITE(6,7)
C      7 FORMAT('0','ANY ADDITIONS?')
C      READ(5,8) I
C      8 FORMAT(A1)
C      IF(I.NE.YES) GO TO 200
C      GO TO 600
C      C CALL THE STANDARD MASS FINDER SUBROUTINE
C      200 CALL STDHNT
C      C LIST THE CURRENT STANDARD MASSES IDENTIFIED BY THE LAST PASS OF THE FINDER ROUTINE
C      300 CALL STDLS
C      C AT THIS POINT THE USER MAY EDIT THE STANDARD MASS LIST
C      400 WRITE(6,9)
C      9 FORMAT('0','ANY CHANGES?')
C      READ(5,8) I
C      KEY=I
C      IF(I.EQ.YES) GO TO 500
C      450 WRITE(6,10)
C      10 FORMAT('0','ANY ADDITIONS?')
C      READ(5,8) I
C      IF(I.EQ.YES) GO TO 600
C      C IF THERE ARE NO CHANGES OR ADDITIONS, THE CURRENT LIST OF STANDARD MASSES IS USED
C      C BY THE DISTANCE TO MASS CONVERSION POLYNOMIAL FOR LEAST SQUARE POLYNOMIAL FITTING.
C      IF(KEY.NE.YES) RETURN
C      GO TO 200
C      500 CALL CHANGE
C      GO TO 450
C      600 CALL ADD
C      GO TO 200
C      END

```

```

C      JANUARY 26,1973                                MICHAEL ROMIEZ
C
C      THIS MODULE WILL SEARCH FOR THE MAKER COMPOUND PEAKS IN ARRAY CENTER
C
C
C
C
C      VARIABLES
C
C      CENTER(1500)  - DISTANCE OF PEAK CENTERS FROM THE LEFT EDGE OF THE PHOTOPLATE
C                    - IN MILLIMETERS
C      HIGHT(1500)   - MAXIMUM INTENSITY VALUE OF EACH PEAK
C      NUMPK          - NUMBER OF PEAKS IN THE PEAK LIST TABLE
C      STDDIS(160)    - THE DISTANCE VALUES OF PEAK CENTERS IDENTIFIED AS STANDARD MASSES
C      STDMA(160)     - THE SQUARE ROOT OF THE MASS VALUE OF IDENTIFIED STANDARD MASSES
C      NUMSTD         - THE NUMBER OF IDENTIFIED STANDARD MASSES
C      REFMA(160)     - A TABLE OF THE MOST INTENSE LINES IN THE MARKER COMPOUND SPECTRUM.
C                    - THESE ARE THE MASSES SEARCHED FOR BY THE STANDARD MASS IDENTIFICATION
C                    - ROUTINE
C      LSTPAR(10)     - PARAMETER LIST FROM PARAMETER LIST MODULE
C      X(200)         - UP TO 200 POINTS THAT ARE USED FOR POLYNOMIAL FITTING
C      Y(200)         - ONE ORDINATE VALUE FOR EACH X VALUE
C      N              - NUMBER OF X AND Y VALUES
C      IDEG           - IT IS THE DEGREE OF THE POLYNOMIAL USED FOR FITTING
C      C(11)          - UP TO ELEVEN COEFFICIENTS OF THE POLYNOMIAL CALCULATED
C      ITOL           - THIS IS A FACTOR USED BY THE STANDARD MASS SEARCHING ROUTINE TO CALCULATE
C                    - THE TOLERANCE WITHIN WHICH IT WILL ACCEPT A PEAK AS A STANDARD MASS. A SMALL
C                    - VALUE MEANS A SMALL TOLERANCE
C      K              - THE REFMA ARRAY POINTER
C      M              - THE CENTER ARRAY POINTER
C      MSTRT          - LOWER LIMIT FOR VALUES OF M
C      MSTOP          - UPPER LIMIT FOR VALUES OF M
C      KSTOP          - UPPER LIMIT FOR VALUES OF K
C      TEMP           - THE DISTANCE WHERE THE NEXT STANDARD MASS IS EXPECTED
C                    - TO BE FOUND
C
C      XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C      SUBROUTINE STDHNT
C      DOUBLE PRECISION CENTER,STDDIS(160),STDMA(160),REFMA(160)
C      DOUBLE PRECISION X(200),Y(200),C(11),TEMP,HTOL,LTOL
C      COMMON/PKLS/ CENTER(1500),HIGHT(1500),NUMPK
C      COMMON/PFKLS/ STDDIS,STDMA,REFMA,NUMSTD
C      COMMON/FIT/ X,Y,NUM,IDEG,C
C      COMMON/PAR/LSTPAR(10)
C
C      SET THE SLIDING TOLERANCE FACTOR
C      ITOL=LSTPAR(9)
C
C      CLEAR VARIABLES
C      DO 100 M=1,200
C      X(M)=0.
C      100 Y(M)=0.
C
C      WE WANT TO USE A QUADRATIC SEARCH ROUTINE FOR WHICH WE NEED
C      THREE PEAKS. THE FIRST TIME WE ENTER THIS SUBROUTINE WE MAY
C      ONLY HAVE TWO IDENTIFIED STANDARD MASSES. A LINEAR SEARCH SECTION
C      IS PROVIDED FOR THIS CASE
C      IF(NUMSTD.GT.2) GO TO 200
C      DO 150 M=1,2
C      X(M)=STDDIS(M)
C      150 Y(M)=STDMA(M)
C      NUM=2
C      IDEG=1
C      CALL POLYFT
C      GO TO 275
C
C      THIS IS THE QUADRATIC SEARCH SECTION
C      200 DO 250 M=1,3
C      X(M)=STDDIS(M)
C      250 Y(M)=STDMA(M)
C      NUM=3
C      IDEG=2
C      CALL POLYFT
C
C      THIS IS THE SEARCHING SECTION
C      275 K=0
C      M=1
C      KSTOP=160
C      MSTOP=NUMPK
C      301 K=K+1
C
C      IF WE HAVE LOOKED FOR ALL THE REFMA ENTRIES, RETURN
C      IF(K.GT.KSTOP) RETURN
C      IF(NUMSTD.LT.3) GO TO 350
C
C      THIS IS THE SECTION WHERE THE EXPECTED DISTANCE VALUE FOR THE NEXT ENTRY
C      IN THE REFMA ARRAY IS CALCULATED.
C      TEMP=(-C(2)+DSQRT(C(2)**2-4*C(3)*(C(1)-REFMA(K)))/(2*C(3))
C      GO TO 351
C      350 TEMP=(REFMA(K)-C(1))/C(2)
C
C      WE MUST SPECIFY A TOLERANCE WITHIN WHICH WE WILL ACCEPT A
C      DISTANCE VALUE AS THE SEARCHED FOR STANDARD MASS. WE HAVE A

```

```

C    JANUARY 26,1973                                MICHAEL ROMIEZ
C
C FORMULA WHICH ALLOWS A HIGHER TOLERANCE AT THE HIGH MASS END. IT
C USES A FACTOR (ITOL) WHICH IS SUPPLIED BY THE PARAMETER
C SETTING MODULE.
351  TOL=(FLOAT(ITOL)*.001/(TEMP/350.))
      HTOL=TEMP+TOL
      LTOL=TEMP-TOL
      GO TO 402
401  M=M+1
C SEARCH ONLY WITHIN THE SUBLIST SPECIFIED BY MSTRT AND MSTOP.
402  IF(M.GT.MSTOP) RETURN
      IF(CENTER(M).LT.LTOL) GO TO 401
      IF(CENTER(M).GT.HTOL) GO TO 301
C WE REACH THIS POINT ONLY IF WE HAVE A HIT.
C IF THE IDENTIFIED STANDARD MASS IS ALREADY IN THE FILE IGNORE
C THIS FIND.
      DO 450 J=1,NUMSTD
        IF(CENTER(M).EQ.STDDIS(J)) GO TO 800
        IF(STDMA(J).EQ.REFMAS(K)) GO TO 800
450  CONTINUE
C IF THE NEW STANDARD MASS IS LARGER THAN THE LARGEST MASS ALREADY
C IN THE FILE THEN WE CAN ADD IT TO THE END OF THE LIST.
500  IF(CENTER(M).LT.STDDIS(NUMSTD)) GO TO 600
      NUMSTD=NUMSTD+1
      STDMA(NUMSTD)=REFMAS(K)
      STDDIS(NUMSTD)=CENTER(M)
      J=NUMSTD
      GO TO 800
C THIS SECTION WILL INSERT AN IDENTIFIED STANDARD MASS WITHIN THE
C STANDARD MASS LIST IN THE PROPER PLACE. THE MASSES MUST BE STORED IN
C ASCENDING ORDER.
600  LIMIT=NUMSTD
      DO 625 J=1,LIMIT
        IF(CENTER(M).GT.STDDIS(J)) GO TO 625
        ITOP=160-J
        DO 650 LL=1,ITOP
          JJ=160-LL
          JJJ=JJ+1
          STDDIS(JJJ)=STDDIS(JJ)
          STDMA(JJJ)=STDMA(JJ)
650  STDDIS(J)=CENTER(M)
          STDMA(J)=REFMAS(K)
          NUMSTD=NUMSTD+1
          GO TO 800
625  CONTINUE
C THIS SECTION CHOOSES THE STANDARD MASSES THAT WILL BE USED TO
C CALCULATE A NEW SET OF COEFFICIENTS. NOTE THAT VARIABLE J
C POINTS TO WHERE IN THE STDDIS LIST THE LAST STANDARD MASS
C WAS INSERTED OR ALREADY FOUND.
800  IF(NUMSTD.EQ.2) GO TO 401
      IF(STDDIS(J+1).NE.0.) J=J+1
      IF(J.LT.3) J=3
      DO 850 NEW=1,3
        INDEX=J-3+NEW
        X(NEW)=STDDIS(INDEX)
850  Y(NEW)=STDMA(INDEX)
      NUM=3
      IDEG=2
      CALL POLYFT
      GO TO 301
      END

```

```

C   JANUARY 29,1973                               MICHAEL ROMIEZ
C
C   THIS MODULE WILL ACCEPT CHANGES AND/OR DELETIONS TO THE STANDARD MASS LIST
C
C                                     VARIABLES
C
C   STDDIS(160)  - THE DISTANCE VALUES OF PEAK CENTERS IDENTIFIED AS STANDARD MASSES
C   STDMA(160)  - THE SQUARE ROOT OF THE MASS VALUE OF IDENTIFIED STANDARD MASSES
C   NUMSTD      - THE NUMBER OF IDENTIFIED STANDARD MASSES
C   REFMAS(160) - A TABLE OF THE MOST INTENSE LINES IN THE MARKER COMPOUND SPECTRUM
C               - THESE ARE THE MASSES SEARCHED FOR BY THE STANDARD MASS IDENTIFICATION
C               - ROUTINE
C
C   XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C   SUBROUTINE CHANGE
C   DOUBLE PRECISION STDDIS(160),STDMA(160),REFMAS(160)
C   COMMON/PFKLST/ STDDIS,STOMA,REFMAS, NUMSTD
5000 WRITE(6,1)
      1 FORMAT('0','ENTER SUBSCRIPT')
      CALL INTGER(MM)
      IF(MM.EQ.0) GO TO 5001
      WRITE(6,3)
      3 FORMAT(1X,'ENTER NEW MASS VALUE')
C IF IT IS DESIRED TO DELETE A PEAK, ENTER ZERO FOR THE MASS VALUE
      READ(5,2) STDMA(MM)
      2 FORMAT(F10.0)
      STDMA(MM)=DSQRT(STDMA(MM))
      GO TO 5000
C AFTER ALL CHANGES AND DELETIONS HAVE BEEN MADE, THE STANDARD MASS LIST MUST BE COMPRESSED.
5001 M=0
5002 M=M+1
5003 IF(STDMA(M).NE.0.) GO TO 5002
      DO 5005 J=M,NUMSTD
      IF(STDMA(J).NE.0.) GO TO 5008
5005 CONTINUE
      NUMSTD=M-1
      RETURN
5008 DO 5010 J=M,NUMSTD
      STDDIS(J)=STDDIS(J+1)
5010 STDMA(J)=STDMA(J+1)
      GO TO 5003
      END

```



```

C      JANUARY 29,1973                                MICHAEL ROMIEZ
C
C      THIS MODULE WILL DISPLAY AREAS OF THE CENTER LIST AND ADD ENTERIES TO THE STANDARD
C      MASS LIST
C
C
C      VARIABLES
C
C      CENTER(1500)      - DISTANCE OF PEAK CENTERS FROM THE LEFT EDGE OF THE PHOTOPLATE
C                        - IN MILLIMETERS
C      HEIGHT(1500)      - MAXIMUM INTENSITY VALUE OF EACH PEAK
C      NUNPK              - NUMBER OF PEAKS IN THE PEAK LIST TABLE
C      STDDIS(160)        - THE DISTANCE VALUES OF PEAK CENTERS IDENTIFIED AS STANDARD MASSES
C      STDMA(160)         - THE SQUARE ROOT OF THE MASS VALUE OF IDENTIFIED STANDARD MASSES
C      NUMSTD             - THE NUMBER OF IDENTIFIED STANDARD MASSES
C      REFMA(160)         - A TABLE OF THE MOST INTENSE LINES IN THE MARKER COMPOUND SPECTRUM.
C                        - THESE ARE THE MASSES SEARCHED FOR BY THE STANDARD MASS IDENTIFICATION
C                        - ROUTINE
C      TEMP               - TEMPORARY STORAGE
C
C      XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C      SUBROUTINE ADD
C      DOUBLE PRECISION STDDIS(160),STDMA(160),REFMA(160),TEMP
C      DOUBLE PRECISION CENTER
C      COMMON/PFKLST/STDDIS,STDMA,REFMA,NUMSTD
C      COMMON/PKLST/CENTER(1500),HIGHT(1500),NUNPK
C      2 WRITE(6,30)
C      30 FORMAT('0','ENTER DISTANCE AREA FOR DISPLAY')
C      ENTER ZERO TO RETURN TO THE CALLING MODULE. ENTER A VALUE LESS THAN 1. TO SKIP THE
C      DISPLAY SECTION AND GO DIRECTLY TO THE PEAK INPUT SECTION.
C      READ(5,31) TEMP
C      31 FORMAT(F10.7)
C      IF(TEMP.EQ.0.) GO TO 999
C      IF(TEMP.LT.1.) GO TO 41
C      M=0
C      1 M=M+1
C      IF(CENTER(M).EQ.0.) GO TO 1000
C      IF(CENTER(M).LT.TEMP) GO TO 1
C      M=M-3
C      K=M+6
C      WRITE(6,32)
C      LIST THE RAW PEAK DATA IN THE VACINITY OF THE DESIRED PEAK
C      32 FORMAT('0',10X,'DISTANCE',10X,'RAW INTENSITY')
C      IF(M.LT.1) M=1
C      DO 40 J=M,K
C      IF(CENTER(J).EQ.0.) GO TO 41
C      40 WRITE(6,33) J,CENTER(J),HIGHT(J)
C      33 FORMAT(4X,13,4X,F8.4,12X,F7.1)
C      41 WRITE(6,34)
C      34 FORMAT('0','ENTER SUBSCRIPT')
C      CALL INTGER(L)
C      IF(L.EQ.0) GO TO 2
C      WRITE(6,35)
C      35 FORMAT(1X,'ENTER MASS')
C      READ(5,36) TEMP
C      36 FORMAT(F10.0)
C      TEMP=DSQRT(TEMP)
C      NN=NUMSTD
C      PLACE THE ADDED PEAK IN THE PROPER PLACE IN THE STANDARD MASS LIST
C      DO 7777 I=1,NN
C      IF(TEMP.GT.STDMA(I)) TO TO 7777
C      LTOP=160-I
C      DO 8000 J=1,LTOP
C      MM=160-J
C      MM=MM+1
C      STDDIS(MMM)=STDDIS(MM)
C      8000 STDMA(MMM)=STDMA(MM)
C      STDDIS(MM)=CENTER(L)
C      STDMA(MM)=TEMP
C      NUMSTD=NUMSTD+1
C      GO TO 2
C      7777 CONTINUE
C      IF THE NEW PEAK IS LARGER THAN THE OTHER ENTERIES, ADD IT TO THE END OF THE LIST.
C      NUMSTD=NUMSTD+1
C      STDDIS(NUMSTD)=CENTER(L)
C      STDMA(NUMSTD)=TEMP
C      GO TO 2
C      1000 WRITE(6,37) CENTER(M-1)
C      37 FORMAT('0','THE LARGEST DISTANCE VALUE IN THE FILE IS',F10.5)
C      GO TO 2
C      999 RETURN
C      END

```

```

C      JANUARY 26,1973                                MICHAEL ROMIEZ
C
C      THIS MODULE WILL LIST THE STANDARD MASSES FOUND BY THE SEARCHING ROUTINE
C
C
C      VARIABLES
C
C      STDDIS(160) - THE DISTANCE VALUES OF PEAK CENTERS IDENTIFIED AS STANDARD MASSES
C      STDMA(160) - THE SQUARE ROOT OF THE MASS VALUE OF IDENTIFIED STANDARD MASSES
C      NUMSTD - THE NUMBER OF IDENTIFIED STANDARD MASSES
C      REFMAS(160) - A TABLE OF THE MOST INTENSE LINES IN THE MARKER COMPOUND SPECTRUM.
C                  - THESE ARE THE MASSES SEARCHED FOR BY THE STANDARD MASS IDENTIFICATION
C                  - ROUTINE
C      X(200) - UP TO 200 ABCISSA VALUES TO BE PASSED TO THE POLYNOMIAL FITTING
C              - ROUTINE
C      Y(200) - UP TO 200 ORDINATE VALUES TO BE PASSED TO THE POLYNOMIAL FITTING
C              - ROUTINE
C      N - NUMBER OF X AND Y VALUES PASSED
C      IDEG - DEGREE OF THE POLYNOMIAL (UP TO TENTH POSSIBLE)
C      C(11) - UP TO 11 COEFFICIENTS RETURNED BY THE POLYNOMIAL ROUTINE
C      LSTPAR(10) - LIST OF PARAMETERS FROM THE PARLIST MODULE
C      PFKMAS(160) - THE MASSES OF IDENTIFIED STANDARD MASSES CALCULATED BY USING THE
C                  - SURROUNDING INUM STANDARD MASSES
C      STD(160) - THE KNOWN EXACT MASS OF EACH IDENTIFIED STANDARD MASS
C      DIFF(160) - THE DIFFERENCE IN MILLIMASS UNITS BETWEEN THE EXACT MASS AND THE
C                  - CALCULATED MASS OF EACH IDENTIFIED STANDARD LINE
C      PPM(16) - THE DIFFERENCE IN PARTS PER MILLION BETWEEN THE EXACT MASS AND THE
C                  - CALCULATED MASS OF EACH IDENTIFIED STANDARD LINE
C      CC(160,11) - THE UP TO 160 SETS OF COEFFICIENTS CALCULATED BY THIS MODULE FOR
C                  - EACH SET OF INUM STANDARD MASSES USED IN THE DISTANCE TO MASS
C                  - CALCULATION
C      INUM - THE NUMBER OF STANDARD MASSES USED FOR EACH SUCCESSIVE
C              - POLYNOMIAL FIT
C      INDEX - POINTER TO THE CC ARRAY
C      IHDEX - THE TOTAL NUMBER OF SETS OF COEFFICIENTS
C      IORDER - THE NUMBER OF COEFFICIENTS RETURNED BY THE POLYNOMIAL FITTING
C              - ROUTINE
C      ISTRT - POINTER TO THE LOWEST OF THE INUM VALUES TO BE TRANSFERED TO
C              - THE POLYNOMIAL FITTING ROUTINE
C      ITOP - POINTER TO THE HIGHEST OF THE INUM VALUES TO BE TRANSFERED TO
C              - THE POLYNOMIAL FITTING ROUTINE
C      M - THE STDDIS AND STDMA ARRAY POINTER
C
C      XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C      SUBROUTINE STLST
C      DOUBLE PRECISION X(200),Y(200),C(11)
C      DOUBLE PRECISION STDMA(160),STDDIS(160),PPM(160),REFMAS(160)
C      DOUBLE PRECISION PFKMAS(160),STD(160),DIFF(160),CC(160,11)
C      COMMON/PFKLST/STDDIS,STDMA,REFMAS, NUMSTD
C      COMMON/FIT/X,Y,N,IDEG,C
C      COMMON/OUTLST/ PFKMAS,STD,DIFF,PPM,CC
C      COMMON/PAR/LSTPAR(10)
C      WRITE(6,11)
C
C      SET THE NUMBER OF STANDARD MASSES USED, AND THE DEGREE OF THE CONVERSION POLYNOMIAL
C      INUM=LSTPAR(7)
C      LIMDEG=LSTPAR(8)
C
C      CLEAR VARIABLES
C      DO 100 M=1,160
C        PFKMAS(M)=0.
C        STD(M)=0.
C        PPM(M)=0.
C        DIFF(M)=0.
C        DO 100 J=1,11
C          C(J)=0.
C        100 CC(M,J)=0.
C
C      INUM IS THE NUMBER OF POINTS USED BY THE DISTANCE TO MASS CONVERSION POLYNOMIAL
C      JNUM IS ONE LESS THAN INUM, AND KNUM IS ONE HALF INUM.
C      JNUM=INUM-1
C      KNUM=INUM/2
C
C      THIS SECTION IS FOR LESS THAN INUM IDENTIFIED STANDARD MASSES
C      IF(NUMSTD.GT.INUM) GO TO 1000
C
C      IF THERE ARE LESS THAN INUM IDENTIFIED STANDARD MASSES, SEND THEM ALL TO CALCULATE
C      A POLYNOMIAL OF DEGREE NUMSTD-2; PROVIDED THAT THE DEGREE DOES NOT EXCEED LIMDEG
C      N=NUMSTD
C      DO 500 M=1,NUMSTD
C        X(M)=STDDIS(M)
C        500 Y(M)=STDMA(M)
C        IDEG=NUMSTD-2
C        IF(IDEG.GT.LIMDEG) IDEG=LIMDEG
C        CALL POLYFT
C        IORDER=IDEG+1
C
C      STORE THE RETURNED SET OF COEFFICIENTS IN THE CC ARRAY; IN THIS CASE THERE WILL BE
C      ONLY ONE SET.
C      DO 501 M=1,IORDER

```

```

C    JANUARY 26,1973                                MICHAEL ROMIEZ
C
501 CC(1,M)=C(M)
    IHDEX=1
    M=1
    GO TO 3000
C THIS SECTION IS FOR MORE THAN INUM IDENTIFIED STANDARD MASSES
1000 INDEX=1
    ISTART=1
1001 ITOP=ISTART+JNUM
    IF(LTOP.GT.NUMSTD) GO TO 2000
    MM=0
C ALL THE STANDARD MASSES ARE SENT TO THE POLYNOMIAL FITTING ROUTINE IN GROUPS OF INUM.
C AFTER ALL THE MASSES HAVE BEEN USED FOR POLYNOMIAL CALCULATION IN THIS MANNER, THERE
C WILL BE NUMSTD-JNUM SETS OF COEFFICIENTS STORED IN THE CC ARRAY. THESE SETS OF COEFFICIENTS
C REPRESENT THE BEST POLYNOMIAL FITS FOR DISTANCE TO MASS CONVERSIONS IN THE NUMSTD-JNUM
C OVERLAPPING AREAS OF THE STANDARD MASS LIST.
    DO 300 M=ISTART,ITOP
        MM=MM+1
        X(MM)=STDDIS(M)
300 Y(MM)=STDMAS(M)
    N=INUM
    IDEG=LIMDEG
C STORE COEFFICIENTS IN THE PROPER PLACE IN THE CC ARRAY
    CALL POLYFT
    DO 350 M=1,11
350 CC(INDEX,M)=C(M)
    INDEX=INDEX+1
    ISTART=ISTART+1
    GO TO 1001
C AFTER ALL SETS OF COEFFICIENTS HAVE BEEN CALCULATED, USE THEM TO DETERMINE THE ERROR
C IN THE DISTANCE TO MASS CONVERSION. LIST THE DIFFERENCE BETWEEN THE KNOWN MASSES OF
C THE REFERENCE PEAKS AND THE MASSES CALCULATED FOR THESE REFERENCE LINES USING THE SAME
C METHOD OF CALCULATION AS FOR UNKNOWN LINES.
2000 INDEX=1
    IHDEX=NUMSTD-JNUM
    M=1
3000 IF(M.GT.NUMSTD) GO TO 5000
    IF(M.LT.KNUM) INDEX=1
    IF(INDEX.GT.IHDEX) INDEX=IHDEX
    PFKMAS(M)=CC(INDEX,1)
    DO 4000 K=1,10
        KK=K+1
4000 PFKMAS(M)=PFKMAS(M)+CC(INDEX,KK)*STDDIS(M)**K
    M=M+1
    INDEX=INDEX+1
    GO TO 3000
5000 DO 4500 K=1,NUMSTD
    PFKMAS(K)=PFKMAS(K)*PFKMAS(K)
    STD(K)=STDMAS(K)*STDMAS(K)
    DIFF(K)=(PFKMAS(K)-STD(K))*1000.
4500 PPM(K)=DABS((DIFF(K)/STD(K))*1000.)
C THIS IS THE LISTING SECTION
    WRITE(6,201)
    WRITE(6,202)
    WRITE(6,203)
    DO 4001 M=1,NUMSTD
4001 WRITE(6,204) M,STDDIS(M),PFKMAS(M),STD(M),DIFF(M),PPM(M)
    RETURN
1 FORMAT('11')
201 FORMAT('10',10X,'DISTANCE',10X,'CALCULATED',10X,'EXACT MASS',10X,'DIFFERENCE',10X,'DIFFERENCE')
202 FORMAT('1',31X,'MASS',35X,'IN MMU',14X,'IN PPM')
203 FORMAT('1')
204 FORMAT(3X,13,5X,F8.4,9X,F10.5,10X,F10.5,13X,F5.2,15X,F5.2)
    END

```



```

C   JANUARY 30,1973                                MICHAEL ROMIEZ
C
C CALL THE ELEMENT FITTING ROUTINE FOR CALCULATION AND DISPLAY ON THE CONVERSATIONAL
C TERMINAL
3000 CALL ELEMFT
      RETURN
100 FORMAT('0','DO YOU WISH TO REVIEW ELEMENT FITTING PARAMETERS?')
102 FORMAT('0','ENTER ISOTOPE NAME')
103 FORMAT('0','IS THIS TERMINAL OUTPUT ?')
104 FORMAT(' ','ENTER MASS')
105 FORMAT(' ','NUMBER FOR THIS ISOTOPE')
106 FORMAT(1X,'THE TOLERANCE FOR ELEMENT FITTING IS',13,' MMU')
51  FORMAT(A1)
52  FORMAT(A4)
53  FORMAT(F10.0)
      END

```

```

C   FEBRUARY 1, 1973                               MICHAEL ROMIEZ
C
C   THIS MODULE WILL CALCULATE AN EXACT MASS FOR EACH UNKNOWN PEAK IN THE PEAK LIST. IT
C   WILL ALSO DETERMINE ALL ELEMENTAL COMPOSITIONS THAT ARE WITHIN A SPECIFIED TOLERANCE
C   OF THE EXACT MASS FROM THE GIVEN LIST OF ELEMENTS.
C
C                                     VARIABLES
C
C   CENTER(1500)  - DISTANCE OF PEAK CENTERS FROM THE LEFT EDGE OF THE PHOTOPLATE
C                 - IN MILLIMETERS
C   HIGHT(1500)  - MAXIMUM INTENSITY VALUE OF EACH PEAK
C   NUMPK        - NUMBER OF PEAKS IN THE PEAK LIST TABLE
C   STDDIS(160)  - THE DISTANCE VALUES OF PEAK CENTERS IDENTIFIED AS STANDARD MASSES
C   STDMA(160)   - THE SQUARE ROOT OF THE MASS VALUE OF IDENTIFIED STANDARD MASSES
C   NUMSTD       - THE NUMBER OF IDENTIFIED STANDARD MASSES
C   REFMA(160)   - A TABLE OF THE MOST INTENSE LINES IN THE MARKER COMPOUND SPECTRUM.
C                 - THESE ARE THE MASSES SEARCHED FOR BY THE STANDARD MASS IDENTIFICATION
C                 - ROUTINE
C   LSTPAR(10)   - LIST OF PARAMETERS FROM THE PARLIST MODULE
C   PEKMA(160)   - THE MASSES OF IDENTIFIED STANDARD MASSES CALCULATED BY USING THE
C                 - SURROUNDING STANDARD MASSES
C   STD(160)     - THE KNOWN EXACT MASS OF EACH IDENTIFIED STANDARD MASS
C   DIFF(160)    - THE DIFFERENCE IN MILLIMASS UNITS BETWEEN THE EXACT MASS AND THE
C                 - CALCULATED MASS OF EACH IDENTIFIED STANDARD LINE
C   PPM(16)      - THE DIFFERENCE IN PARTS PER MILLION BETWEEN THE EXACT MASS AND THE
C                 - CALCULATED MASS OF EACH IDENTIFIED STANDARD LINE
C   CC(160,11)   - THE UP TO 160 SETS OF COEFFICIENTS CALCULATED FOR EACH SET OF
C                 - STANDARD MASSES USED IN THE DISTANCE TO MASS CALCULATION
C   MASS(12)     - UP TO TWELVE MASSES OF ELEMENTS
C   NAME(12)     - A FOUR CHARACTER NAME FOR UP TO TWELVE ELEMENTS
C   NUMBER(12)   - THE MAXIMUM NUMBER OF EACH ELEMENT USED IN ELEMENT FITTING
C   NTOL         - THE TOLERANCE IN MILLIMASS UNITS USED FOR ELEMENT FITTING
C   NUM          - THE NUMBER OF ELEMENTS IN THE MASS ARRAY
C   CALMA(391)   - A LIST OF THE MASSES OF THE PEAKS IN THE MARKER COMPOUND THAT
C                 - WILL BE REMOVED FROM THE CLCMA TABLE IF FOUND
C   CLCMA(1500)  - A CALCULATED MASS FOR EACH PEAK IN ARRAY CENTER
C   ATNUM(12)    - THE NUMBER OF ATOMS OF EACH ELEMENT THAT FITS A PARTICULAR
C                 - ELEMENTAL COMPOSITION
C   COUNT1      - A COUNTER THAT CONTROLS THE OUTPUT LISTING
C   COUNT2      - THE NUMBER OF ELEMENTAL COMPOSITIONS FOUND SO FAR FOR THE
C                 - THE PARTICULAR CLCMA VALUE UNDER CONSIDERATION
C
C   XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C
C   SUBROUTINE ELEMFT
C   DOUBLE PRECISION CENTER,STDMA(160),STDDIS(160),PPM(160),REFMA(160)
C   DOUBLE PRECISION PEKMA(160),STD(160),DIFF(160),CC(160,11)
C   DOUBLE PRECISION CLCMA(1500),DIF,HTOL,LTOL,CALMA(391),MASS
C   INTEGER COUNT1,COUNT2,ATNUM(12)
C   REAL MAX
C   COMMON/PFKLST,STDDIS,STDMA,REFMA,NUMSTD
C   COMMON/PAR/ LSTPAR(10)
C   COMMON/OUTLST/ PEKMA,STD,DIFF,PPM,CC
C   COMMON/PKLST/ CENTER(1500),HIGHT(1500),NUMPK
C   COMMON/ELEM/ MASS(12),NAME(12),NUMBER(12),NTOL,NUM
C   COMMON/COMP/CLCMA,ATNUM,COUNT1,COUNT2,DIF,IMAS
C
C   CLEAR CLCMA ARRAY
C   DO 50 I=1,1500
C   50 CLCMA(I)=0.
C
C   SET THE NUMBER OF STANDARD MASSES USED FOR POLYNOMIAL FITTING IN MODULE STDST
C   INUM=LSTPAR(7)
C
C   CALL THE MARKER COMPOUND LISTING MODULE
C   CALL STDST
C
C   INITIALIZE THE COUNTER THAT CONTROLS THE OUTPUT FORMAT
C   COUNT1=45
C
C   CALCULATE THE EXACT MASS OF EACH PEAK USING THE PROPER SET OF POLYNOMIAL COEFFICIENTS
C   FROM MODULE STDST. INDEX IS THE CC ARRAY POINTER, IHDEX IS THE POINTER TO THE LAST
C   SET OF COEFFICIENTS IN THE CC ARRAY.
C   1012 IHDEX=1
C   IHDEX=NUMSTD-INUM+1
C   IF (IHDEX.LT.1) IHDEX=1
C   M=1
C   1 DO 1000 J=1,NUMSTD
C   IF (STDDIS(J).LT.CENTER(M)) GO TO 1000
C   GO TO 1001
C   1000 CONTINUE
C   1001 INDEX=J-(INUM/2)
C   IF (J.LE.INUM/2) INDEX=1
C   IF (INDEX.GT.IHDEX) INDEX=IHDEX
C   CLCMA(M)=CC (INDEX,1)
C   DO 1005 K=1,10
C   KK=K+1
C   1005 CLCMA(M)=CLCMA(M)+CC (INDEX, KK)*CENTER(M)**K

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C FEBRUARY 1, 1973

MICHAEL ROMIEZ

```
C
M=M+1
IF(M.GT.NUMPK) GO TO 2000
GO TO 1
2000 DO 1010 M=1,NUMPK
1010 CLCMAS(M)=CLCMAS(M)*CLCMAS(M)
C READ IN MASSES TO BE STRIPED OUT
READ(4) CALMAS
C STRIP OUT BACKGROUND AND MARKER COMPOUND PEAKS
TOL=FLOAT(NTOL)*.001
J=0
10 J=J+1
IF(J.GT.NUMPK) GO TO 2500
DO 2001 M=1,391
HTOL=CALMAS(M)+TOL
LTOL=CALMAS(M)-TOL
IF(CLCMAS(J).LT.LTOL) GO TO 10
IF(CLCMAS(J).GT.HTOL) GO TO 2001
CLCMAS(J)=0.
GO TO 10
2001 CONTINUE
GO TO 10
C COMPRESS CLCMAS ARRAY
2500 M=0
100 M=M+1
170 IF(CLCMAS(M).NE.0.) GO TO 100
DO 150 J=M,NUMPK
CLCMAS(J)=CLCMAS(J+1)
CENTER(J)=CENTER(J+1)
150 HIGHT(J)=HIGHT(J+1)
DO 167 K=M,NUMPK
IF(CLCMAS(K).EQ.0.) GO TO 167
GO TO 170
167 CONTINUE
C IF THERE ARE NO PEAKS IN THE MASS SPECTRUM, RETURN TO THE CALLING MODULE
IF(CLCMAS(1).EQ.0) RETURN
C CALCULATE NEW NUMPK VALUE
2700 DO 2800 L=1,NUMPK
IF(CLCMAS(L).NE.0.) GO TO 2800
LL=L-1
TO TO 2900
2800 CONTINUE
2900 NUMPK=LL
C NORMALIZE INTENSITY VALUES
MAX=HIGHT(1)
DO 4950 J=2,NUMPK
IF(HIGHT(J).GT.MAX) MAX=HIGHT(J)
4950 CONTINUE
IF(MAX.EQ.0) GO TO 3005
DO 4951 J=1,NUMPK
4951 HIGHT(J)=(HIGHT(J)/MAX)*100.
C CALCULATE ELEMENTAL COMPOSITIONS
3005 NUM3=NUMBER(3)
NUM4=NUMBER(4)
NUM5=NUMBER(5)
NUM6=NUMBER(6)+1
NUM7=NUMBER(7)+1
NUM8=NUMBER(8)+1
NUM9=NUMBER(9)+1
NUM10=NUMBER(10)+1
NUM11=NUMBER(11)+1
NUM12=NUMBER(12)+1
DO 4000 M=1,NUMPK
HTOL=CLCMAS(M)+TOL
LTOL=CLCMAS(M)-TOL
COUNT2=0
NUM2=1DINT(HTOL/12.)+1
NUM1=NUM2*2+8
DO 4050 I12=1,NUM12
I112=I12-1
ATNUM(12)=I112
DO 4050 I11=1,NUM11
I111=I11-1
ATNUM(11)=I111
DO 4050 I10=1,NUM10
I110=I10-1
ATNUM(10)=I110
DO 4050 I9=1,NUM9
I19=I9-1
ATNUM(9)=I19
DO 4050 I8=1,NUM8
I18=I8-1
```


C FEBRUARY 1, 1973

MICHAEL ROMIEZ

C

```
ATNUM(8)=118
DO 4050 17=1,NUM7
117=17-1
ATNUM(7)=117
DO 4050 16=1,NUM6
116=16-1
ATNUM(6)=116
DO 4050 15=1,NUM5
115=15-1
ATNUM(5)=115
DO 4050 14=1,NUM4
114=14-1
ATNUM(4)=114
DO 4050 13=1,NUM3
113=13-1
ATNUM(3)=113
DO 4050 12=1,NUM2
112=12-1
ATNUM(2)=112
SUM=0.
DO 4100 11=2,NUM
```

4100 SUM=SUM+ATNUM(11)*MASS(11)

C ITEST IS AN INTEGER VALUE WHICH IS USED TO FIT THE ELEMENT HYDROGEN INTO THE

C ELEMENTAL COMPOSITIONS. IF ITEST IS LESS THAN ZERO THIS MEANS THAT SUM

C IS GREATER THAN CLCMAS(M) PLUS THE TOLERANCE AND WE WILL DISMISS THIS COMPOSITION.

C WHEN ITEST IS ZERO OR POSITIVE WE HAVE A POSSIBLE COMPOSITION AND WE MAKE FURTHER

C TESTS

```
ITEST=IDINT(HTOL-SUM)
IF(ITEST.GT.NUM1) GO TO 4050
IF(ITEST) 4050,4051,4051
```

```
4051 ATNUM(1)=ITEST
SUM=SUM+MASS(1)*ATNUM(1)
GO TO 4053
```

C WE CONTINUE TO ADD HYDROGEN ATOMS TO THE COMBINATIONS OF OTHER ELEMENTS UNTIL WE HAVE

C A FIT OR WE HAVE TOO HIGH OF A TOTAL MASS FOR A FIT.

```
4052 SUM=SUM+MASS(1)
ATNUM(1)=ATNUM(1)+1
4053 IF(SUM.GT.HTOL) GO TO 4050
IF SUM.LT.LTOL) GO TO 4052
COUNT2=COUNT2+1
DIF=(CLCMAS(M)-SUM)*1000.
IMAS=M
```

CALL OUTPUT

4050 CONTINUE

C IF THERE WERE NO ELEMENTAL COMPOSITIONS FOUND FOR THIS MASS, LIST THIS INFORMATION

C AS PART OF THE OUTPUT.

```
IMAS=M
IF(COUNT2.EQ.0) CALL OUTPUT
```

```
4000 CONTINUE
RETURN
END
```

```

C   FEBRUARY 1,1973                               MICHAEL ROMIEZ
C
C   THIS MODULE LISTS ELEMENTAL COMPOSITIONS FOUND BY THE ELEMENT FITTING ROUTINE
C
C                                     VARIABLES
C
C   CENTER(1500)      - DISTANCE OF PEAK CENTERS FROM THE LEFT EDGE OF THE PHOTOPLATE
C                     - IN MILLIMETERS
C   HIGHT(1500)       - MAXIMUM INTENSITY VALUE OF EACH PEAK
C   NMPK              - NUMBER OF PEAKS IN THE PEAK LIST TABLE
C   MASS(12)          - UP TO TWELVE MASSES OF ELEMENTS
C   NAME(12)          - A FOUR CHARACTER NAME FOR UP TO TWELVE ELEMENTS
C   NUMBER(12)        - THE MAXIMUM NUMBER OF EACH ELEMENT USED IN ELEMENT FITTING
C   NTOL              - THE TOLERANCE IN MILLIMASS UNITS USED FOR ELEMENT FITTING
C   NUM               - THE NUMBER OF ELEMENTS IN THE MASS ARRAY
C   ATNUM(12)         - THE NUMBER OF ATOMS OF EACH ELEMENT THAT FITS A PARTICULAR
C                     - ELEMENTAL COMPOSITION
C   COUNT1            - A COUNTER THAT CONTROLS THE OUTPUT LISTING
C   COUNT2            - THE NUMBER OF ELEMENTAL COMPOSITIONS FOUND SO FAR FOR THE
C                     - THE PARTICULAR CLCMAS VALUE UNDER CONSIDERATION
C
C   XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
C   SUBROUTINE OUTPUT
C   DOUBLE PRECISION CENTER,MASS,CLCMAS(1500),DIFF
C   INTEGER ATNUM(12),COUNT1,COUNT2
C   COMMON/ELEM/MASS(12),NAME(12),NUMBER(12),NTOL,NUM
C   COMMON/PKLIST/ CENTER(1500),HIGHT(1500),NMPK
C   COMMON/COMP/CLCMAS,ATNUM,COUNT1,COUNT2,DIFF,IMAS
C TEST NEED FOR A NEW PAGE HEADING
C   IF(COUNT1.GE.45) GO TO 1002
C IF COUNT2 IS ZERO, THERE HAVE BEEN NO ELEMENTAL FITS FOR THIS MASS
500  IF(COUNT2.EQ.0) GO TO 1001
C IF THIS IS A NEW MASS WE SKIP A SPACE
C   IF(COUNT2.EQ.1) GO TO 1003
1000 WRITE(6,303) CENTER(IMAS),HIGHT(IMAS),CLCMAS(IMAS),DIFF,(ATNUM(M),M=1,NUM)
C   COUNT1=COUNT1+1
C   RETURN
1001 WRITE(6,304) CENTER(IMAS),HIGHT(IMAS),CLCMAS(IMAS),NTOL
C   COUNT1=COUNT1+2
C   RETURN
1002 WRITE(6,100)
C   WRITE(6,300)
C   WRITE(6,301) (NAME(M),M=1,NUM)
C   WRITE(6,302)
C   COUNT1=5
C   GO TO 500
1003 WRITE(6,203)
C   COUNT1=COUNT1+1
C   GO TO 1000
203  FORMAT(' ')
100  FORMAT(' ')
300  FORMAT(6X,'DISTANCE',6X,'RELATIVE',7X,'MASS',7X,'DIFFERENCE')
301  FORMAT(20X,'INTENSITY',19X,'IN MMU',5X,12A5)
302  FORMAT('0')
303  FORMAT(6X,F8.4,7X,F5.1,5X,F10.5,8X,F5.2,3X,12I5)
304  FORMAT('0',5X,F8.4,7X,F5.1,5X,F10.5,10X,'NO ELEMENTAL COMPOSITIONS FOUND WITHIN',13,' MMU')
C   END

```

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POSTMASTER :

If Undeliverable (Section 158
Postal Manual) Do Not Return

"The aeronautical and space activities of the United States shall be conducted so as to contribute . . . to the expansion of human knowledge of phenomena in the atmosphere and space. The Administration shall provide for the widest practicable and appropriate dissemination of information concerning its activities and the results thereof."

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